

GenCore version 5.1.6  
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OM protein - protein search, using sw model

Run on: December 3, 2003, 11:48:05 ; Search time 26.3333 Seconds  
(without alignments)  
58.797 Million cell updates/sec

Title: US-09-912-414-2  
Perfect score: 45  
Sequence: 1 WVRWHF 6

Scoring table: BLOSUM62  
Gapop 10.0 , Gapext 0.5

Searched: 830525 seqs, 258052604 residues  
Total number of hits satisfying chosen parameters: 3526

Minimum DB seq length: 0  
Maximum DB seq length: 15

Post-processing: Minimum Match 0%  
Maximum Match 100%  
Listing first 45 summaries

- Database : SPTREMBL.23:\*
- 1: sp\_archea:\*
  - 2: sp\_bacteria:\*
  - 3: sp\_fungi:\*
  - 4: sp\_human:\*
  - 5: sp\_invertebrate:\*
  - 6: sp\_mammal:\*
  - 7: sp\_mhc:\*
  - 8: sp\_organelle:\*
  - 9: sp\_phage:\*
  - 10: sp\_plant:\*
  - 11: sp\_rodent:\*
  - 12: sp\_virus:\*
  - 13: sp\_vertebrate:\*
  - 14: sp\_unclassified:\*
  - 15: sp\_rvrius:\*
  - 16: sp\_bacteriap:\*
  - 17: sp\_archheap:\*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	DB ID	Description
1	28	62.2	9	8 Q8SHF0	Q8shf0 chamaeleo n
2	24	53.3	8	8 Q94VF6	Q94vf6 varanus job
3	24	53.3	10	8 P92632	P92632 eremias gra
4	23	51.1	10	8 Q9TG41	Q9tg41 ophisaurus
5	22	48.9	8	8 Q94VJ4	Q94vj4 varanus ben
6	22	48.9	8	13 P79940	P79940 xenopus lae
7	22	48.9	10	8 Q9B4X0	Q9b4x0 notophthalm
8	22	48.9	10	8 Q958L2	Q958l2 rana tempor
9	22	48.9	10	8 Q958L8	Q958l8 rana catesb
10	22	48.9	10	8 Q958K6	Q958k6 rana pretio
11	22	48.9	10	8 Q958K0	Q958k0 rana cascad
12	22	48.9	10	8 Q958L5	Q958l5 rana sylvat
13	22	48.9	10	8 Q958K3	Q958k3 rana aurora
14	22	48.9	10	8 Q94NH4	Q94nh4 rana muscos
15	22	48.9	10	8 Q94VD2	Q94vd2 varanus pan
16	21	46.7	8	8 Q94VC1	Q94vc1 varanus rud

17	21	46.7	8	8 Q9TD02	Q9td02 terranatos
18	21	46.7	8	8 Q9T4Y2	Q9t4y2 asterina pe
19	21	46.7	9	8 Q9T688	Q9t688 gecko gecko
20	21	46.7	10	8 Q9T8K7	Q9t8k7 liolaemus m
21	21	46.7	10	8 Q9T8N1	Q9t8n1 liolaemus p
22	21	46.7	10	8 Q79903	Q79903 oplurus cuv
23	21	46.7	10	8 Q8W969	Q8w969 anolis orto
24	21	46.7	10	8 Q8WDH8	Q8wdh8 anolis mest
25	21	46.7	10	8 Q9T8T6	Q9t8t6 liolaemus m
26	21	46.7	10	8 Q9T8L3	Q9t8l3 liolaemus l
27	21	46.7	10	8 P92616	P92616 aspidosceli
28	21	46.7	10	8 Q9T8G8	Q9t8g8 liolaemus c
29	21	46.7	10	8 Q958K9	Q958k9 rana boylli
30	21	46.7	10	8 Q9TFU9	Q9tfu9 teratoscinc
31	21	46.7	10	8 Q9T8X7	Q9t8x7 phymaturus
32	21	46.7	10	8 Q79885	Q79885 anolis pate
33	21	46.7	10	8 Q9T8Q5	Q9t8q5 liolaemus l
34	21	46.7	10	8 P92654	P92654 euprepis au
35	21	46.7	10	8 Q9T8L0	Q9t8l0 liolaemus o
36	21	46.7	10	8 Q9T8W8	Q9t8w8 liolaemus b
37	21	46.7	10	8 Q9T8R4	Q9t8r4 liolaemus p
38	21	46.7	10	8 Q9T8M8	Q9t8m8 liolaemus m
39	21	46.7	10	8 Q9T8S1	Q9t8s1 liolaemus l
40	21	46.7	10	8 Q9T8S4	Q9t8s4 liolaemus c
41	21	46.7	10	8 Q9ZYU4	Q9zyu4 sceloporos
42	21	46.7	10	8 P92758	P92758 teratoscinc
43	21	46.7	10	8 Q9T8T9	Q9t8t9 liolaemus l
44	21	46.7	10	8 Q9ZYT5	Q9zyt5 uta stansbu
45	21	46.7	10	8 Q9T8J8	Q9t8j8 liolaemus w

ALIGNMENTS

RESULT 1  
Q8SHF0  
ID Q8SHF0 PRELIMINARY; PRT; 9 AA.  
AC Q8SHF0;  
DT 01-JUN-2002 (TrEMBLrel. 21, Created)  
DT 01-JUN-2002 (TrEMBLrel. 21, Last sequence update)  
DT 01-JUN-2002 (TrEMBLrel. 21, Last annotation update)  
DE Cytochrome c oxidase subunit I (Fragment).  
GN COI.  
OS Chamaeleo namaquensis.  
OG Mitochondrion.  
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
OC Lepidosauria; Squamata; Iguania; Acrodonta; Chamaeleonidae; Chamaeleo.  
OX NCBI\_TaxID=179917;  
RN [1]  
RP SEQUENCE FROM N.A.  
RA Townsend T.M., Larson A.L.;  
RT "Molecular Phylogenetics and Mitochondrial Genomic Evolution in the  
Chamaeleonidae (Reptilia, Squamata).";  
RL Submitted (NOV-2001) to the EMBL/GenBank/DBJ databases.  
DR EMBL; AF448757; AAL90553.1; -.  
KW Mitochondrion.  
FT NON TER 9  
SQ SEQUENCE 9 AA; 1205 MW; 358CB72733640733 CRC64;

Query Match 62.2%; Score 28; DB 8; Length 9;  
Best Local Similarity 75.0%; Pred. No. 8.3e+05;  
Matches 3; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

Qy 1 WVRW 4  
|:|:  
Db 2 WLRW 5

RESULT 2  
Q94VF6  
ID Q94VF6 PRELIMINARY; PRT; 8 AA.  
AC Q94VF6;  
DT 01-DEC-2001 (TrEMBLrel. 19, Created)

DT 01-DEC-2001 (TReMBLrel. 19, Last sequence update)  
DT 01-DEC-2001 (TReMBLrel. 19, Last annotation update)  
DE Cytochrome c oxidase subunit I (Fragment).  
GN COI.  
OS Varanus jobiensis.  
OG Mitochondrion.  
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
OC Lepidosauria; Squamata; Scleroglossa; Anguimorpha; Varanidae; Varanus.  
OX NCBI\_TaxID=169843;  
RN [1]  
RP SEQUENCE FROM N.A.  
RA Ast J.C.;  
RT "Mitochondrial DNA evidence and evolution in Varanoidea (Squamata).";  
RL Cladistics 17:0-0(2001).  
DR EMBL; AF407507; AAL10075.1; -.  
KW Mitochondrion.  
FT NON\_TER 8  
SQ SEQUENCE 8 AA; 1144 MW; EFD729DB436411A6 CRC64;

Query Match 53.3%; Score 24; DB 8; Length 8;  
Best Local Similarity 75.0%; Pred. No. 8.3e+05;  
Matches 3; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

QY 3 RWHF 6  
|||  
Db 3 RWYF 6

RESULT 3  
P92632  
ID P92632; PRELIMINARY; PRT; 10 AA.  
AC P92632;  
DT 01-MAY-1997 (TReMBLrel. 03, Created)  
DT 01-MAY-1997 (TReMBLrel. 03, Last sequence update)  
DT 01-NOV-1998 (TReMBLrel. 08, Last annotation update)  
DE Cytochrome c oxidase subunit I (Fragment).  
GN COI.  
OS Eremias grammica.  
OG Mitochondrion.  
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
OC Lepidosauria; Squamata; Scleroglossa; Scincormorpha; Lacertoidea;  
OC Lacertidae; Eremias.  
OX NCBI\_TaxID=52179;  
RN [1]  
RP SEQUENCE FROM N.A.  
RX MEDLINE=97153826; PubMed=9000757;  
RA Macey J.R., Larson A., Ananjeva N.B., Fang Z., Papenfuss T.J.;  
RT "Two novel gene orders and the role of light-strand replication in  
rearrangement of the vertebrate mitochondrial genome.";  
RL Mol. Biol. Evol. 14:91-104(1997).  
RN [2]  
RP SEQUENCE FROM N.A.  
RX MEDLINE=97153820; PubMed=9000751;  
RA Macey J.R., Larson A., Ananjeva N.B., Papenfuss T.J.;  
RT "Replication slippage may cause parallel evolution in the secondary  
structures of mitochondrial transfer RNAs.";  
RL Mol. Biol. Evol. 14:30-39(1997).  
DR EMBL; U71331; AAB48277.1; -.  
KW Mitochondrion.  
FT NON\_TER 10  
SQ SEQUENCE 10 AA; 1288 MW; 5B3580C9D3640057 CRC64;

Query Match 53.3%; Score 24; DB 8; Length 10;  
Best Local Similarity 60.0%; Pred. No. 7.5e+02;  
Matches 3; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

QY 2 VRWHF 6  
:|  
Db 4 IRWFF 8

RESULT 4  
Q9TG41

ID Q9TG41  
AC Q9TG41; PRELIMINARY; PRT; 10 AA.  
DT 01-MAY-2000 (TReMBLrel. 13, Created)  
DT 01-MAY-2000 (TReMBLrel. 13, Last sequence update)  
DT 01-MAY-2000 (TReMBLrel. 13, Last annotation update)  
DE Cytochrome c oxidase subunit I (Fragment).  
GN COI.  
OS Ophisaurus apodus.  
OG Mitochondrion.  
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
OC Lepidosauria; Squamata; Scleroglossa; Anguimorpha; Anguidae;  
OC Ophisaurus.  
OX NCBI\_TaxID=102191;  
RN [1]  
RP SEQUENCE FROM N.A.  
RX MEDLINE=99343613; PubMed=104113621;  
RA Macey J.R., Schulte J.A. II, Larson A., Tuniyev B.S., Orlov N.,  
RA Papenfuss T.J.;  
RT "Molecular phylogenetics, tRNA evolution, and historical biogeography  
in anguillid lizards and related taxonomic families.";  
RL Mol. Phylogenet. Evol. 12:250-272(1999).  
DR EMBL; AF085623; AAD51559.1; -.  
KW Mitochondrion.  
FT NON\_TER 10  
SQ SEQUENCE 10 AA; 1239 MW; 1A3580C7336412C0 CRC64;

Query Match 51.1%; Score 23; DB 8; Length 10;  
Best Local Similarity 80.0%; Pred. No. 1.1e+03;  
Matches 4; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

QY 2 VRWHF 6  
|||  
Db 4 VRWLF 8

RESULT 5

Q94VJ4  
ID Q94VJ4 PRELIMINARY; PRT; 8 AA.  
AC Q94VJ4;  
DT 01-DEC-2001 (TReMBLrel. 19, Created)  
DT 01-DEC-2001 (TReMBLrel. 19, Last sequence update)  
DT 01-DEC-2001 (TReMBLrel. 19, Last annotation update)  
DE Cytochrome c oxidase subunit I (Fragment).  
GN COI.  
OS Varanus bengalensis nebulosis.  
OG Mitochondrion.  
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
OC Lepidosauria; Squamata; Scleroglossa; Anguimorpha; Varanidae; Varanus.  
OX NCBI\_TaxID=169827;  
RN [1]  
RP SEQUENCE FROM N.A.  
RA Ast J.C.;  
RT "Mitochondrial DNA evidence and evolution in Varanoidea (Squamata).";  
RL Cladistics 17:0-0(2001).  
DR EMBL; AF407492; AAL10031.1; -.  
KW Mitochondrion.  
FT NON\_TER 8  
SQ SEQUENCE 8 AA; 1053 MW; E8B5B9C733640056 CRC64;

Query Match 48.9%; Score 22; DB 8; Length 8;  
Best Local Similarity 60.0%; Pred. No. 8.3e+05;  
Matches 3; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

QY 2 VRWHF 6  
:|  
Db 2 IRWLF 6

RESULT 6  
P79940  
ID P79940 PRELIMINARY; PRT; 8 AA.  
AC P79940;  
DT 01-MAY-1997 (TReMBLrel. 03, Created)

DT 01-MAY-1997 (TrEMBLrel. 03, Last sequence update)  
DT 01-DEC-2001 (TrEMBLrel. 19, Last annotation update)  
DE xMeisl-4 protein (Fragment).  
OS Xenopus laevis (African clawed frog).  
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
OC Amphibia; Batrachia; Anura; Mesobatrachia; Pipidae;  
OC Xenopodinae; Xenopus.  
OX NCBI\_TaxID=8355;  
RN [1]  
RP SEQUENCE FROM N.A.  
RX MEDLINE=97202105; PubMed=9049632;  
RA Steelman S., Moscow J.J., Muzynski K., North C., Druck T.,  
RA Montgomery J.C., Huebner K., Daar I.O., Buchberg A.M.;  
RT "Identification of a conserved family of Meisl-related homeobox  
genes.";  
RL Genome Res. 7:142-156(1997).  
DR EMBL; U68389; AAB19199.1; -.  
DR TRANSPAC; T03410; -.  
FT NON TER 1  
SQ SEQUENCE 8 AA; 1187 MW; 278B51F37B11F40B CRC64;  
  
Query Match 48.9%; Score 22; DB 13; Length 8;  
Best Local Similarity 66.7%; Pred. No. 8.3e+05;  
Matches 2; Conservative 1; Mismatches 0; Indels 0; Gaps 0;  
  
QY 4 WHF 6  
Db 5 WHY 7  
  
RESULT 7  
Q9B4X0  
ID Q9B4X0 PRELIMINARY; PRT; 10 AA.  
AC Q9B4X0;  
DT 01-JUN-2001 (TrEMBLrel. 17, Created)  
DT 01-JUN-2001 (TrEMBLrel. 17, Last sequence update)  
DT 01-JUN-2001 (TrEMBLrel. 17, Last annotation update)  
DE Cytochrome c oxidase subunit 1 (Fragment).  
GN COI.  
OS Notophthalmus viridescens (Eastern newt) (Triturus viridescens).  
OG Mitochondrion.  
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
OC Amphibia; Batrachia; Caudata; Salamandroidea; Salamandridae;  
OC Notophthalmus.  
OX NCBI\_TaxID=8316;  
RN [1]  
RP SEQUENCE FROM N.A.  
RX MEDLINE=21175761; PubMed=11277635;  
RA Weisrock D.W., Macey J.R., Ugurtas I.H., Larson A., Papenfuss T.J.;  
RT "Molecular Phylogenetics and Historical Biogeography among  
Salamanders of the 'True' Salamander Clade: Rapid Branching of  
RT Numerous Highly Divergent Lineages in Mertensiella luschani Associated  
RT with the Rise of Anatolia.";  
RL Mol. Phylogenet. Evol. 18:434-448(2001).  
DR EMBL; AF296616; AAK30305.1; -.  
KW Mitochondrion.  
FT NON TER 10  
SQ SEQUENCE 10 AA; 1298 MW; 03D380C733640050 CRC64;  
  
Query Match 48.9%; Score 22; DB 8; Length 10;  
Best Local Similarity 60.0%; Pred. No. 1.5e+03;  
Matches 3; Conservative 1; Mismatches 1; Indels 0; Gaps 0;  
  
QY 2 VRWHF 6  
Db 4 IRWLF 8  
  
RESULT 8  
Q958L2  
ID Q958L2 PRELIMINARY; PRT; 10 AA.  
AC Q958L2;  
DT 01-DEC-2001 (TrEMBLrel. 19, Created)

DT 01-DEC-2001 (TrEMBLrel. 19, Last sequence update)  
DT 01-DEC-2001 (TrEMBLrel. 19, Last annotation update)  
DE Cytochrome c oxidase subunit I (Fragment).  
GN COI.  
OS Rana temporaria (European common frog).  
OG Mitochondrion.  
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
OC Amphibia; Batrachia; Anura; Neobatrachia; Ranioidea; Rana.  
OX NCBI\_TaxID=8407;  
RN [1]  
RP SEQUENCE FROM N.A.  
RX MEDLINE=21184280; PubMed=11286498;  
RA Macey J.R., Strasburg J.L., Brisson J.A., Vredenburg V.T.,  
RA Jennings M., Larson A.;  
RT "Molecular Phylogenetics of Western North American Frogs of the Rana  
RT boylei Species Group.";  
RL Mol. Phylogenet. Evol. 19:131-143(2001).  
DR EMBL; AF314018; AAK56874.1; -.  
KW Mitochondrion.  
FT NON TER 10  
SQ SEQUENCE 10 AA; 1354 MW; C0D380C9D36411A9 CRC64;  
  
Query Match 48.9%; Score 22; DB 8; Length 10;  
Best Local Similarity 50.0%; Pred. No. 1.5e+03;  
Matches 3; Conservative 1; Mismatches 2; Indels 0; Gaps 0;  
  
QY 1 WVRWHF 6  
Db 3 FTRWFF 8  
  
RESULT 9  
Q958L8  
ID Q958L8 PRELIMINARY; PRT; 10 AA.  
AC Q958L8;  
DT 01-DEC-2001 (TrEMBLrel. 19, Created)  
DT 01-DEC-2001 (TrEMBLrel. 19, Last sequence update)  
DT 01-DEC-2001 (TrEMBLrel. 19, Last annotation update)  
DE Cytochrome c oxidase subunit I (Fragment).  
GN COI.  
OS Rana catesbeiana (Bull frog).  
OG Mitochondrion.  
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
OC Amphibia; Batrachia; Anura; Neobatrachia; Ranioidea; Rana.  
OX NCBI\_TaxID=8400;  
RN [1]  
RP SEQUENCE FROM N.A.  
RX MEDLINE=21184280; PubMed=11286498;  
RA Macey J.R., Strasburg J.L., Brisson J.A., Vredenburg V.T.,  
RA Jennings M., Larson A.;  
RT "Molecular Phylogenetics of Western North American Frogs of the Rana  
RT boylei Species Group.";  
RL Mol. Phylogenet. Evol. 19:131-143(2001).  
DR EMBL; AF314016; AAK56868.1; -.  
KW Mitochondrion.  
FT NON TER 10  
SQ SEQUENCE 10 AA; 1354 MW; C0D380C9D36411A9 CRC64;  
  
Query Match 48.9%; Score 22; DB 8; Length 10;  
Best Local Similarity 50.0%; Pred. No. 1.5e+03;  
Matches 3; Conservative 1; Mismatches 2; Indels 0; Gaps 0;  
  
QY 1 WVRWHF 6  
Db 3 FTRWFF 8  
  
RESULT 10  
Q958K6  
ID Q958K6 PRELIMINARY; PRT; 10 AA.  
AC Q958K6;  
DT 01-DEC-2001 (TrEMBLrel. 19, Created)  
DT 01-DEC-2001 (TrEMBLrel. 19, Last sequence update)

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DT 01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
DE Cytochrome c oxidase subunit I (Fragment).
GN COI.
OS Rana pretiosa.
OG Mitochondrion.
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Amphibia; Batrachia; Anura; Neobatrachia; Ranoidea; Ranidae; Rana.
OX NCBI_TaxID=69834;
RN [1]
RP SEQUENCE FROM N.A.
RX MEDLINE=21184280; PubMed=11286498;
RA Macey J.R., Strasburg J.L., Brisson J.A., Vredenburg V.T.,
RA Jennings M., Larson A.;
RT "Molecular Phylogenetics of Western North American Frogs of the Rana
RT boyllii Species Group.";
RL Mol. Phylogenet. Evol. 19:131-143(2001).
DR EMBL; AF314020; AAK56880.1; -.
KW Mitochondrion.
FT NON TER 10
SQ SEQUENCE 10 AA; 1354 MW; COD380C9D36411A9 CRC64;

Query Match 48.9%; Score 22; DB 8; Length 10;
Best Local Similarity 50.0%; Pred. No. 1.5e+03;
Matches 3; Conservative 1; Mismatches 2; Indels 0; Gaps 0;

QY 1 WVRWHF 6
Db : |||
3 FTRWFF 8

RESULT 11
Q958K0 PRELIMINARY; PRT; 10 AA.
AC Q958K0;
DT 01-DEC-2001 (TrEMBLrel. 19, Created)
DT 01-DEC-2001 (TrEMBLrel. 19, Last sequence update)
DT 01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
DE Cytochrome c oxidase subunit I (Fragment).
GN COI.
OS Rana cascadae.
OG Mitochondrion.
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Amphibia; Batrachia; Anura; Neobatrachia; Ranoidea; Ranidae; Rana.
OX NCBI_TaxID=160497;
RN [1]
RP SEQUENCE FROM N.A.
RX MEDLINE=21184280; PubMed=11286498;
RA Macey J.R., Strasburg J.L., Brisson J.A., Vredenburg V.T.,
RA Jennings M., Larson A.;
RT "Molecular Phylogenetics of Western North American Frogs of the Rana
RT boyllii Species Group.";
RL Mol. Phylogenet. Evol. 19:131-143(2001).
DR EMBL; AF314022; AAK56886.1; -.
KW Mitochondrion.
FT NON TER 10
SQ SEQUENCE 10 AA; 1354 MW; COD380C9D36411A9 CRC64;

Query Match 48.9%; Score 22; DB 8; Length 10;
Best Local Similarity 50.0%; Pred. No. 1.5e+03;
Matches 3; Conservative 1; Mismatches 2; Indels 0; Gaps 0;

QY 1 WVRWHF 6
Db : |||
3 FTRWFF 8

RESULT 12
Q958L5 PRELIMINARY; PRT; 10 AA.
AC Q958L5;
DT 01-DEC-2001 (TrEMBLrel. 19, Created)
DT 01-DEC-2001 (TrEMBLrel. 19, Last sequence update)
DT 01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
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DE Cytochrome c oxidase subunit I (Fragment).
GN COI.
OS Rana sylvatica (Wood frog).
OG Mitochondrion.
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Amphibia; Batrachia; Anura; Neobatrachia; Ranoidea; Ranidae; Rana.
OX NCBI_TaxID=45438;
RN [1]
RP SEQUENCE FROM N.A.
RX MEDLINE=21184280; PubMed=11286498;
RA Macey J.R., Strasburg J.L., Brisson J.A., Vredenburg V.T.,
RA Jennings M., Larson A.;
RT "Molecular Phylogenetics of Western North American Frogs of the Rana
RT boyllii Species Group.";
RL Mol. Phylogenet. Evol. 19:131-143(2001).
DR EMBL; AF314017; AAK56871.1; -.
KW Mitochondrion.
FT NON TER 10
SQ SEQUENCE 10 AA; 1354 MW; COD380C9D36411A9 CRC64;

Query Match 48.9%; Score 22; DB 8; Length 10;
Best Local Similarity 50.0%; Pred. No. 1.5e+03;
Matches 3; Conservative 1; Mismatches 2; Indels 0; Gaps 0;

QY 1 WVRWHF 6
Db : |||
3 FTRWFF 8

RESULT 13
Q958K3 PRELIMINARY; PRT; 10 AA.
AC Q958K3;
DT 01-DEC-2001 (TrEMBLrel. 19, Created)
DT 01-DEC-2001 (TrEMBLrel. 19, Last sequence update)
DT 01-MAR-2003 (TrEMBLrel. 23, Last annotation update)
DE Cytochrome c oxidase subunit I (Fragment).
GN COI.
OS Rana aurora (Red-legged frog).
OG Mitochondrion.
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Amphibia; Batrachia; Anura; Neobatrachia; Ranoidea; Ranidae; Rana.
OX NCBI_TaxID=160496;
RN [1]
RP SEQUENCE FROM N.A.
RX MEDLINE=21184280; PubMed=11286498;
RA Macey J.R., Strasburg J.L., Brisson J.A., Vredenburg V.T.,
RA Jennings M., Larson A.;
RT "Molecular Phylogenetics of Western North American Frogs of the Rana
RT boyllii Species Group.";
RL Mol. Phylogenet. Evol. 19:131-143(2001).
DR EMBL; AF314021; AAK56883.1; -.
KW Mitochondrion.
FT NON TER 10
SQ SEQUENCE 10 AA; 1354 MW; COD380C9D36411A9 CRC64;

Query Match 48.9%; Score 22; DB 8; Length 10;
Best Local Similarity 50.0%; Pred. No. 1.5e+03;
Matches 3; Conservative 1; Mismatches 2; Indels 0; Gaps 0;

QY 1 WVRWHF 6
Db : |||
3 FTRWFF 8

RESULT 14
Q94NH4 PRELIMINARY; PRT; 10 AA.
AC Q94NH4;
DT 01-DEC-2001 (TrEMBLrel. 19, Created)
DT 01-DEC-2001 (TrEMBLrel. 19, Last sequence update)
DT 01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
DE Cytochrome c oxidase subunit I (Fragment).
```



XX OS Synthetic.  
XX PN WO200044771-A1.  
XX XX  
PD 03-AUG-2000.  
XX XX  
PF 26-JAN-2000; 2000WO-GB00227.  
XX XX  
PR 26-JAN-1999; 99GB-0001710.  
XX PA (PROL-) PROLIFIX LTD.  
XX PI Mueller R, Kontermann RE, Montigiani S;  
XX XX  
DR WPI; 2000-532806/48.  
XX XX  
PT Peptides binding to the DNA binding domain of transcription factor E2F  
PT and inhibiting cell cycle progression, useful for the treatment of  
PT cancer  
XX XX  
PS Example; Page 26; 42pp; English.  
XX XX  
CC Peptides which bind to the DNA binding domain of transcription  
CC factor E2F and inhibit cell cycle progression may be useful as  
CC research agents to investigate the interaction between E2F and DP-1,  
CC or the activation of transcription by E2F-1/DP-1 heterodimers. They  
CC may also be used for inducing apoptosis and/or cell cycle arrest in  
CC a cell, particularly for treatment of cancer or other proliferative  
CC disorders such as psoriasis and restenosis.  
XX XX  
SQ Sequence 6 AA;  
Query Match 86.7%; Score 39; DB 21; Length 6;  
Best Local Similarity 100.0%; Pred. No. 9.3e+05;  
Matches 5; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  
QY 1 WVRWH 5  
Db 1 WVRWH 5  
RESULT 5  
AAB01508  
ID AAB01508 standard; peptide; 6 AA.  
XX AC AAB01508;  
XX DT 08-NOV-2000 (first entry)  
XX DE Peptide which binds to transcription factor E2F-1 DNA binding domain.  
KW DNA binding; transcription factor; E2F; E2F-1; cell cycle; DP-1;  
KW activation; transcription; apoptosis; proliferative disorder;  
KW psoriasis; restenosis.  
XX OS Synthetic.  
XX PN WO200044771-A1.  
XX XX  
PD 03-AUG-2000.  
XX XX  
PF 26-JAN-2000; 2000WO-GB00227.  
XX XX  
PR 26-JAN-1999; 99GB-0001710.  
XX PA (PROL-) PROLIFIX LTD.  
XX PI Mueller R, Kontermann RE, Montigiani S;  
XX XX  
DR WPI; 2000-532806/48.  
XX XX  
PT Peptides binding to the DNA binding domain of transcription factor E2F  
PT and inhibiting cell cycle progression, useful for the treatment of  
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PT and inhibiting cell cycle progression, useful for the treatment of  
PT cancer  
XX XX  
PS Example; Page 26; 42pp; English.  
XX XX  
CC Peptides which bind to the DNA binding domain of transcription  
CC factor E2F and inhibit cell cycle progression may be useful as  
CC research agents to investigate the interaction between E2F and DP-1,  
CC or the activation of transcription by E2F-1/DP-1 heterodimers. They  
CC may also be used for inducing apoptosis and/or cell cycle arrest in  
CC a cell, particularly for treatment of cancer or other proliferative  
CC disorders such as psoriasis and restenosis.  
XX XX  
SQ Sequence 6 AA;  
Query Match 77.8%; Score 35; DB 21; Length 6;  
Best Local Similarity 83.3%; Pred. No. 9.3e+05;  
Matches 5; Conservative 0; Mismatches 1; Indels 0; Gaps 0;  
QY 1 WVRWHF 6  
Db 1 WVRWAF 6  
RESULT 6  
AAB01499  
ID AAB01499 standard; peptide; 6 AA.  
XX AC AAB01499;  
XX DT 08-NOV-2000 (first entry)  
XX DE Peptide which binds to transcription factor E2F-1 DNA binding domain.  
KW DNA binding; transcription factor; E2F; E2F-1; cell cycle; DP-1;  
KW activation; transcription; apoptosis; proliferative disorder;  
KW psoriasis; restenosis.  
XX OS Synthetic.  
XX PN WO200044771-A1.  
XX XX  
PD 03-AUG-2000.  
XX XX  
PF 26-JAN-2000; 2000WO-GB00227.  
XX XX  
PR 26-JAN-1999; 99GB-0001710.  
XX PA (PROL-) PROLIFIX LTD.  
XX PI Mueller R, Kontermann RE, Montigiani S;  
XX XX  
DR WPI; 2000-532806/48.  
XX XX  
PT Peptides binding to the DNA binding domain of transcription factor E2F  
PT and inhibiting cell cycle progression, useful for the treatment of  
PT cancer  
XX XX  
PS Claim 4; Page 9; 42pp; English.  
XX XX  
CC Peptides which bind to the DNA binding domain of transcription  
CC factor E2F and inhibit cell cycle progression may be useful as  
CC research agents to investigate the interaction between E2F and DP-1,  
CC or the activation of transcription by E2F-1/DP-1 heterodimers. They  
CC may also be used for inducing apoptosis and/or cell cycle arrest in  
CC a cell, particularly for treatment of cancer or other proliferative  
CC disorders such as psoriasis and restenosis.

XX SQ Sequence 6 AA;  
Query Match 75.6%; Score 34; DB 21; Length 6;  
Best Local Similarity 66.7%; Pred. No. 9.3e+05;  
Matches 4; Conservative 0; Mismatches 2; Indels 0; Gaps 0;  
Qy 1 WVRWHF 6  
| | | |  
Db 1 WXXWHF 6  
RESULT 7  
AAB01504  
ID AAB01504 standard; peptide; 6 AA.  
XX AC AAB01504;  
XX DT 08-NOV-2000 (first entry)  
XX DE Peptide which binds to transcription factor E2F-1 DNA binding domain.  
XX KW DNA binding; transcription factor; E2F; E2F-1; cell cycle; DP-1;  
XX KW activation; transcription; apoptosis; proliferative disorder;  
XX KW psoriasis; restenosis.  
XX OS Synthetic.  
XX PN WO200044771-A1.  
XX PD 03-AUG-2000.  
XX PF 26-JAN-2000; 2000WO-GB00227.  
XX PR 26-JAN-1999; 99GB-0001710.  
XX PA (PROL-) PROLIFIX LTD.  
XX PI Mueller R, Kontermann RE, Montigiani S;  
XX DR WPI; 2000-532806/48.  
XX PT Peptides binding to the DNA binding domain of transcription factor E2F  
XX PT and inhibiting cell cycle progression, useful for the treatment of  
XX PT cancer  
XX PS Example; Page 26; 42pp; English.  
XX CC Peptides which bind to the DNA binding domain of transcription  
XX CC factor E2F and inhibit cell cycle progression may be useful as  
XX CC research agents to investigate the interaction between E2F and DP-1,  
XX CC or the activation of transcription by E2F-1/DP-1 heterodimers. They  
XX CC may also be used for inducing apoptosis and/or cell cycle arrest in  
XX CC a cell, particularly for treatment of cancer or other proliferative  
XX CC disorders such as psoriasis and restenosis.  
XX SQ Sequence 6 AA;  
Query Match 75.6%; Score 34; DB 21; Length 6;  
Best Local Similarity 100.0%; Pred. No. 9.3e+05;  
Matches 5; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  
Qy 2 VRWHF 6  
| | | |  
Db 2 VRWHF 6  
RESULT 8  
AAR60429  
ID AAR60429 standard; peptide; 8 AA.  
XX AC AAR60429;  
XX DE Antiproliferative peptide to transplantable human B-cell lymphoma.

DT 25-MAR-2003 (updated)  
DT 30-MAR-1995 (first entry)  
XX DE Antiproliferative peptide to transplantable human B-cell lymphoma.  
XX KW antiproliferative; transplant; B-cell lymphoma line SUP-B8; Burkitt's;  
XX KW inhibit clonal expansion; induce apoptosis; anti-idiotypic; Igm lambda;  
XX KW inhibit cell proliferation; peptidomimetics; cell surface receptor;  
XX KW immunoglobulin superfamily; treatment; neoplasia; identification;  
XX KW induce replication; therapy; clonal anergy; modulate tyrosine kinase.  
XX OS Synthetic.  
XX PN WO9418345-A1.  
XX PD 18-AUG-1994.  
XX PF 04-FEB-1994; 94WO-US01319.  
XX PR 05-FEB-1993; 93US-0014426.  
XX PR 15-NOV-1993; 93US-0153341.  
XX PA (AFFY-) AFFYMAX TECHNOLOGIES NV.  
XX PA (STRD ) UNIV LELAND STANFORD JUNIOR.  
XX PI Bhatt RR, Dower WJ, Levy R, Renschler MF;  
XX DR WPI; 1994-279762/34.  
XX PT Identifying anti-proliferative peptide(s) which specifically bind  
XX PT to immunoglobulin super-family species idiotype - esp. to inhibit  
XX PT B-cell lymphoma and leukocytic leukaemia cell proliferation, for  
XX PT anti-idiotypic therapy  
XX PS Claim 7; Page 45; 69pp; English.  
XX CC AAR60400-73 are peptide ligands which bind to purified Igm lambda  
XX CC receptor of the human Burkitt's lymphoma cell line SUP-B8. Peptides  
XX CC AAR60414 to AAR60473 were biotinylated and linked to streptavidin.  
XX CC The peptides were identified with the use of filamentous phage  
XX CC libraries displaying random peptides. Corresponding synthetic  
XX CC peptides bound specifically to this Ig receptor, and blocked the  
XX CC binding of an anti-idiotypic antibody. The ligands, when conjugated  
XX CC to form dimers or tetramers, induced cell death by apoptosis in  
XX CC vitro at nanomolar concentrations. This effect was associated with  
XX CC the specific stimulation of intracellular protein tyrosine  
XX CC phosphorylation. The peptides of the invention can be used individually,  
XX CC as complexes of cross-linked peptides or can be conjugated to deliver  
XX CC toxins or radionuclides to neoplastic cells bearing the specific Ig  
XX CC receptor.  
XX CC (Updated on 25-MAR-2003 to correct PN field.)  
XX SQ Sequence 8 AA;  
Query Match 75.6%; Score 34; DB 15; Length 8;  
Best Local Similarity 80.0%; Pred. No. 9.3e+05;  
Matches 4; Conservative 0; Mismatches 1; Indels 0; Gaps 0;  
Qy 1 WVRWH 5  
| | | |  
Db 3 WYRWH 7  
RESULT 9  
AAR60444  
ID AAR60444 standard; peptide; 8 AA.  
XX AC AAR60444;  
XX DT 25-MAR-2003 (updated)  
XX DT 30-MAR-1995 (first entry)  
XX DE Antiproliferative peptide to transplantable human B-cell lymphoma.

XX KW antiproliferative; transplant; B-cell lymphoma line SUP-B8; Burkitt's;  
KW inhibit clonal expansion; induce apoptosis; anti-idiotype; IgM lambda;  
KW inhibit cell proliferation; peptidomimetics; cell surface receptor;  
KW immunoglobulin superfamily; treatment; neoplasia; identification;  
KW induce replication; therapy; clonal anergy; modulate tyrosine kinase.  
XX OS Synthetic.  
XX WO9418345-A1.  
XX 18-AUG-1994.  
XX 04-FEB-1994; 94WO-US01319.  
XX 05-FEB-1993; 93US-0014426.  
XX 15-NOV-1993; 93US-0153341.  
XX (AFFY-) AFFYMAX TECHNOLOGIES NV.  
XX (STRD ) UNIV LELAND STANFORD JUNIOR.  
XX Bhatt RR, Dower WJ, Levy R, Renschler MF;  
XX WPI; 1994-279762/34.  
XX Identifying anti-proliferative peptide(s) which specifically bind  
XX to immunoglobulin super-family species idiotype - esp. to inhibit  
XX B-cell lymphoma and leukocytic leukaemia cell proliferation, for  
XX anti-idiotype therapy  
XX Claim 7; Page 45; 69pp; English.  
XX AAR60400-73 are peptide ligands which bind to purified IgM lambda  
XX receptor of the human Burkitt's lymphoma cell line SUP-B8. Peptides  
XX AAR60414 to AAR60473 were biotinylated and linked to streptavidin.  
XX The peptides were identified with the use of filamentous phage  
XX libraries displaying random peptides. Corresponding synthetic  
XX peptides bound specifically to this Ig receptor, and blocked the  
XX binding of an anti-idiotype antibody. The ligands, when conjugated  
XX to form dimers or tetramers, induced cell death by apoptosis in  
XX vitro at nanomolar concentrations. This effect was associated with  
XX the specific stimulation of intracellular protein tyrosine  
XX phosphorylation. The peptides of the invention can be used individually,  
XX as complexes of cross-linked peptides or can be conjugated to deliver  
XX toxins or radionuclides to neoplastic cells bearing the specific Ig  
XX receptor.  
XX (Updated on 25-MAR-2003 to correct PN field.)  
XX SQ Sequence 8 AA;  
Query Match 75.6%; Score 34; DB 15; Length 8;  
Best Local Similarity 80.0%; Pred. No. 9.3e+05;  
Matches 4; Conservative 0; Mismatches 1; Indels 0; Gaps 0;  
QY 1 WVRWH 5  
Db 3 WYRWH 7  
RESULT 10  
AAR37389  
ID AAR37389 standard; peptide; 6 AA.  
XX AAR37389;  
AC AAR37389;  
XX 07-JUL-1993 (first entry)  
DT Peptide for treating septic shock.  
XX Toxic shock; blood endotoxin removal; serum; diagnostic reagent;  
KW cytokine release control; treatment; pertussis; bacterial meningitis;  
KW HIV related infections; polymyxin B; Group II.  
XX

OS Synthetic.  
XX Key Location/Qualifiers  
FH Region 1..3  
FT /note= "repeat region"  
FT Region 4..6  
FT /note= "repeat region"  
XX ZA9200943-A.  
XX 25-NOV-1992.  
XX 10-FEB-1992; 92ZA-0000943.  
XX 11-FEB-1991; 91US-0658744.  
XX (PORR/) PORRO M.  
XX Porro M;  
XX WPI; 1993-094304/11.  
XX New peptide for treatment or prevention of toxic shock - comprises  
XX specified sequences of aminoacid(s) and analogs  
XX comprising sequences retro-orientated  
XX Example; Page 5; 39pp; English.  
XX The (Group II) peptide is an example of a generic peptide of formula  
XX R-( Lys/Arg/His - Phe/Tyr/Trp - Leu/Ile/Val )n-R, where n = 1-100  
XX and each R is H, an amino acid residue or a fatty acid residue.  
XX The peptide is useful for treating or preventing septic shock,  
XX mixing with polymyxin B to reduce its toxicity; removing  
XX endotoxins from blood, sera or other fluids (in vivo or in  
XX vitro); controlling release of cytokines induced by endotoxins;  
XX as diagnostic reagents to detect and quantify toxins in blood  
XX or sera; preparing non-toxic antigenic complexes of lipid A or  
XX lipopolysaccharide (LPS); and for treating pertussis, bacterial  
XX meningitis and HIV-related infections. The usual dose is 10-100  
XX ug/kg/day, given parenterally. It binds to the same sites as  
XX polymyxin B, i.e. it inhibits all the toxic effects of lipid A. It  
XX has no antibiotic activity; does not lyse erythrocytes; has no  
XX toxicity in mice when injected at 50mg/kg and is relatively unstable  
XX against proteases.  
XX SQ Sequence 6 AA;  
Query Match 68.9%; Score 31; DB 14; Length 6;  
Best Local Similarity 100.0%; Pred. No. 9.3e+05;  
Matches 4; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  
QY 1 WVRW 4  
Db 2 WVRW 5  
RESULT 11  
AAW28912  
ID AAW28912 standard; peptide; 6 AA.  
XX AAW28912;  
AC AAW28912;  
XX 20-JAN-1998 (first entry)  
DT Opioid peptide.  
XX enkephalin; mu-opioid receptor ligand; agonist; antagonist.  
XX Synthetic.  
XX Key Location/Qualifiers  
FH Modified-site 1  
FT /note= "N-acetyl-Arg"  
FT

FT Modified-site 6 /note= "the C-terminal is in amide form"  
FT  
XX US5641861-A.  
PN  
XX 24-JUN-1997.  
PD  
XX 07-JUN-1995; 95US-0487006.  
PF  
XX 07-JUN-1995; 95US-0487006.  
PR  
XX (TORR-) TORREY PINES INST MOLECULAR STUDIES.  
PA  
XX Dooley CT, Houghten RA;  
PI  
XX WPI; 1997-340994/31.  
DR  
XX New opioid peptides which bind mu receptors specifically - have  
PT agonist or antagonist activity and are used for study and  
PT localisation of mu receptors and to treat peripheral side effects of  
PT morphine etc.  
XX  
PS Disclosure; Column 8; 92pp; English.  
XX  
CC The patent discloses the following new peptides, which are opioids which  
CC bind specifically to the mu receptor: Ac-Phe-Arg-Trp-Trp-Tyr-Xaa-NH2 (1);  
CC Ac-Arg-Trp-Ile-Gly-Trp-Xaa-NH2 (2); Trp-Trp-Pro-Lys-His-Xaa-NH2 (3);  
CC Trp-Trp-Pro-Xaa1-NH2 (4); Tyr-Pro-Phe-Gly-Phe-Xaa-NH2 (5);  
CC D-Ile-D-Met-D-Ser-D-Trp-D-Trp-(Gly)n-Xaa2-NH2 (6);  
CC D-Ile-D-Met-D-Thr-D-Trp-Gly-Xaa2-NH2 (7); Tyr-Al-B2-C3-NH2 (214);  
CC Pm and red (Me)x(H)y-Tyr-(NMe)z-Tyr-(Xaa3)z-NH2 (221); and  
CC Trp-Trp-Pro-D4-(His)z-(Xaa)z-NH2 (222); where Xaa = any natural amino  
CC acid; Xaa1 = Lys or Arg; n and z = 0 or 1; Xaa2 = Gly or the D form of  
CC any naturally occurring amino acid; Al = D-norvaline or D-norleucine;  
CC B2 = Gly, Phe or Trp; C3 = Trp or naphthylalanine; x and y = 0-2, but  
CC not over 2 in total; Xaa3 = Phe, DPhe or benzylamino; D4 = Lys or Arg;  
CC Pm and red indicate permethylation and reduction of all CO in peptide  
CC links to methylene. These new compounds are useful: (i) for in vitro  
CC assay and study of opiate receptor subtypes, particularly mu receptors  
CC in the brain; (ii) for in vivo localisation of receptor subtypes; and  
CC (iii) therapeutically to block the peripheral effects (e.g. constipation  
CC and pruritus) of centrally acting pain killers such as morphine.  
CC They are very selective for the mu opioid receptor, over binding to the  
CC delta and kappa receptor subtypes.  
CC The present sequence is a specific example of a peptide (2).  
XX  
SQ Sequence 6 AA;  
Query Match 68.9%; Score 31; DB 18; Length 6;  
Best Local Similarity 60.0%; Pred. No. 9.3e+05;  
Matches 3; Conservative 1; Mismatches 1; Indels 0; Gaps 0;  
QY 1 WVRWH 5  
Db 2 WIGWH 6  
RESULT 12  
AAR93770  
ID AAR93770 standard; Protein; 6 AA.  
XX  
AC AAR93770;  
XX  
XX 23-SEP-1997 (first entry)  
DT  
XX New peptide which acts as mu-opioid receptor ligand.  
DE  
XX mu-receptor; opioid; opiate; agonist; antagonist; diagnosis;  
KW analgesic.  
XX Synthetic.  
OS  
XX Key Location/Qualifiers  
FH

FT Modified-site 1 /note= "N-acetyl-Arg"  
FT Misc-difference 6 /note= "this residue is in C-terminal amide form"  
FT  
XX WO9640208-A1.  
PN  
XX 19-DEC-1996.  
PD  
XX 06-JUN-1996; 96WO-US09321.  
PF  
XX 07-JUN-1995; 95US-0476438.  
PR  
XX (TORR-) TORREY PINES INST MOLECULAR STUDIES.  
PA  
XX Dooley CT, Houghten RA;  
PI  
XX WPI; 1997-051895/05.  
DR  
XX New mu opioid receptor binding ligand peptide(s) - useful for  
PT in-vitro and in-vivo diagnosis, as analgesics, and for blocking  
PT peripheral effects of centrally acting drugs, e.g. morphine  
XX  
PS Disclosure; Page 19; 57pp; English.  
XX  
CC The patent discloses eight new groups of opioid peptides which bind  
CC to the mu-receptor to act as agonists or antagonists. The peptides  
CC can be used for in-vitro assays to study opiate receptor subtypes  
CC (especially the mu type) in brain or other tissue samples; and for  
CC in-vivo diagnosis to localise opioid subtypes. The peptides are also  
CC useful as drugs to treat pathologies associated with other compounds  
CC which interact with the opioid receptor system. Therefore they can be  
CC used in medicaments for treating pathologies associated with the mu  
CC receptor and as analgesics. They can be used therapeutically to block  
CC the peripheral effects of centrally acting pain killers, e.g. to  
CC prevent side effects such as constipation and pruritis associated  
CC with morphine. The present sequence represents a specific example  
CC of one of the new groups of peptides, of formula  
CC Ac-Arg-Trp-Ile-Gly-Trp-Xaa-NH2 where Xaa = a naturally occurring  
CC amino acid.  
XX  
SQ Sequence 6 AA;  
Query Match 68.9%; Score 31; DB 18; Length 6;  
Best Local Similarity 60.0%; Pred. No. 9.3e+05;  
Matches 3; Conservative 1; Mismatches 1; Indels 0; Gaps 0;  
QY 1 WVRWH 5  
Db 2 WIGWH 6  
RESULT 13  
AAY23019  
ID AAY23019 standard; peptide; 6 AA.  
XX  
AC AAY23019;  
XX  
XX 23-AUG-1999 (first entry)  
DT  
XX Opioid peptide which inhibits binding of enkephalin.  
DE  
XX Opioid peptide; ligand binding; opioid receptor;  
KW micro-selective opioid peptide; enkephalin; opioid receptor system;  
KW blocking; peripheral effect; centrally acting pain killer; morphine.  
XX Synthetic.  
OS  
XX Key Location/Qualifiers  
FH Modified-site 1 /note= "acetylated"  
FT Modified-site 6 /note= "amidated"  
FT

XX US5919897-A.  
PN 06-JUL-1999.  
XX  
PF 07-JUN-1995; 95US-0488659.  
XX  
PR 07-JUN-1995; 95US-0488659.  
XX  
PA (TORR-) TORREY PINES INST MOLECULAR STUDIES.  
XX  
PI Dooley CT, Houghten RA;  
XX  
DR WPI; 1999-394647/33.  
XX  
PT New opioid peptides useful for blocking the peripheral effects of  
PT centrally acting pain killers such as morphine  
XX  
PS Example 1; Column 8; 92pp; English.  
XX  
CC The specification describes opioid peptides, in which each of the  
CC N atoms in the peptide backbone between respective amino acids is  
CC modified by permethylation, perallylation, perethylation, perbenzylation  
CC and pernapthylation. The peptides inhibit ligand binding to an opioid  
CC receptor. Specifically, the peptides inhibit the micro-selective  
CC opioid peptide enkephalin. The peptides can be used in vivo  
CC diagnostically to localize opioid receptor subtypes. They can be used  
CC to treat pathologies associated with other compounds which interact with  
CC the opioid receptor system. The peptides are especially useful for  
CC blocking the peripheral effects of centrally acting pain killers such  
CC as morphine. AAY23005-Y23024 represent opioid peptides of the invention,  
CC and are derived from the general sequence given in AAY23004.  
XX  
SQ Sequence 6 AA;  
Query Match 68.9%; Score 31; DB 20; Length 6;  
Best Local Similarity 60.0%; Pred. No. 9.3e+05;  
Matches 3; Conservative 1; Mismatches 1; Indels 0; Gaps 0;  
QY 1 WVRWH 5  
|: ||  
Db 2 WIGWH 6  
RESULT 14  
AAB01507  
ID AAB01507 standard; peptide; 6 AA.  
XX  
AC AAB01507;  
XX  
DT 08-NOV-2000 (first entry)  
XX  
DE Peptide which binds to transcription factor E2F-1 DNA binding domain.  
XX  
KW DNA binding; transcription factor; E2F; E2F-1; cell cycle; DP-1;  
KW activation; transcription; apoptosis; proliferative disorder;  
KW psoriasis; restenosis.  
XX  
OS Synthetic.  
XX  
PN WO200044771-A1.  
XX  
PD 03-AUG-2000.  
XX  
PF 26-JAN-2000; 2000WO-GB00227.  
XX  
PR 26-JAN-1999; 99GB-0001710.  
XX  
PA (PROL-) PROLIFIX LTD.  
XX  
PI Mueller R, Kontermann RE, Montigiani S;  
XX  
DR WPI; 2000-532806/48.

XX Peptides binding to the DNA binding domain of transcription factor E2F  
PT and inhibiting cell cycle progression, useful for the treatment of  
PT cancer  
XX  
PS Example; Page 26; 42pp; English.  
XX  
CC Peptides which bind to the DNA binding domain of transcription  
CC factor E2F and inhibit cell cycle progression may be useful as  
CC research agents to investigate the interaction between E2F and DP-1,  
CC or the activation of transcription by E2F-1/DP-1 heterodimers. They  
CC may also be used for inducing apoptosis and/or cell cycle arrest in  
CC a cell, particularly for treatment of cancer or other proliferative  
CC disorders such as psoriasis and restenosis.  
XX  
SQ Sequence 6 AA;  
Query Match 68.9%; Score 31; DB 21; Length 6;  
Best Local Similarity 83.3%; Pred. No. 9.3e+05;  
Matches 5; Conservative 0; Mismatches 1; Indels 0; Gaps 0;  
QY 1 WVRWHF 6  
||| ||  
Db 1 WVRAHF 6  
RESULT 15  
AAM45777  
ID AAM45777 standard; Peptide; 7 AA.  
XX  
AC AAM45777;  
XX  
DT 25-OCT-2001 (first entry)  
XX  
DE H11 binding site consensus conforming peptide (CCP) #2048.  
XX  
KW Antigen-binding; tumour; diagnosis; stress protein-peptide complex; SPPC;  
KW immunogenically cross-reactive; cancer; immunogenic cancer cell;  
KW cytostatic; vaccine; tumour-specific immunogenic response inducer;  
KW astrocytoma; fibrosarcoma; myxosarcoma; liposarcoma; oligodendroglioma;  
KW ependymoma; medulloblastoma; primitive neural ectodermal tumour.  
XX  
OS Homo sapiens.  
OS Synthetic.  
XX  
PN CA2290722-A1.  
XX  
PD 08-JUN-2001.  
XX  
PF 08-DEC-1999; 99CA-2290722.  
XX  
PR 08-DEC-1999; 99CA-2290722.  
XX  
PA (NOVO-) NOVOPHARM BIOTECH INC.  
XX  
PI Kaplan HA, Maiti PK, Fast DG, Herman W, Dan MD, Lewis KE;  
PI Entwistle JM, MacDonald GC;  
XX  
DR WPI; 2001-425937/46.  
XX  
PT Composition useful for treating and diagnosing cancer, comprises stress  
PT protein-peptide complexes associated with tumor, and isolated  
PT antigen-binding fragments of an antibody that binds specifically to the  
PT complex  
XX  
PS Example 4; Page 108; 154pp; English.  
XX  
CC The present invention describes a composition (I) comprising stress  
CC protein-peptide complexes (SPPC) associated with tumours that is  
CC specifically immunogenically cross-reactive with cell surface-associated  
CC SPSCs specific to target cancer (TC). Also described is an isolated  
CC antigen-binding fragment of an antibody that binds specifically to SPSCs  
CC or a population of different SPSCs consisting of immunogenic cancer cell



CC surface-associated SPPC of TC. (I) has cytostatic activity and can be  
 CC used in vaccine production and as a tumour-specific immunogenic response  
 CC inducer. (I) is useful for treating 71 types of cancers or tumours in a  
 CC subject, such as astrocytoma, fibrosarcoma, myxosarcoma, liposarcoma,  
 CC oligodendroglioma, ependymoma, medulloblastoma, and primitive neural  
 CC ectodermal tumour (PNET). (I) is useful as cancer immunogen including  
 CC vaccines. (I) is useful for diagnostic and palliative use, for detecting  
 CC or imaging cancer cells, and to monitor the course of amelioration of  
 CC malignancy in an individual. AAM43707 to AAM47109 represent peptides  
 CC which are used in the exemplification of the present invention.

XX  
 SQ Sequence 7 AA;

Query Match 68.9%; Score 31; DB 22; Length 7;  
 Best Local Similarity 66.7%; Pred. No. 9.3e+05;  
 Matches 4; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

QY 1 WVRWHF 6  
 | | | | |  
 DB 1 WRRWNF 6

RESULT 16  
 AAR86140  
 ID AAR86140 standard; peptide; 10 AA.  
 AC AAR86140;  
 XX  
 DT 26-JUN-1996 (first entry)  
 XX  
 DE Anti-ELAM-1 binding peptide #117.  
 XX  
 KW Peptide mimetic; endothelial leukocyte adhesion molecule; ELAM; selectin;  
 KW receptor; leukocyte; vascular wall; endothelium; extravasation;  
 KW inflammation; sialyl Lewis; cell surface glycoprotein; HL60 cell.  
 XX  
 OS Synthetic.  
 XX  
 PN WO9531210-A1.  
 PD 23-NOV-1995.  
 XX  
 PF 11-MAY-1995; 95WO-US06315.  
 XX  
 PR 11-MAY-1994; 94US-0241054.  
 XX  
 PA (AFFY-) AFFYMAX TECHNOLOGIES NV.  
 XX  
 PI Barrett RW, Cwirla SE, Dower WJ, Koller KJ, Lee J;  
 PI Martens CL, Ruhland-fritsch B;  
 XX  
 DR WPI; 1996-010687/01.  
 XX  
 PT New peptide(s) that bind to endothelial leukocyte adhesion molecule  
 PT 1 - useful for treating inflammation and other E-selectin mediated  
 PT diseases  
 XX  
 PS Disclosure; Page 17; 85pp; English.  
 XX  
 CC Peptides AAR86024-R86236 are examples of peptides and their mimetics  
 CC that bind to endothelial leukocyte adhesion molecule (ELAM)-1. This  
 CC molecule is a member of the selectin family of receptors and is involved  
 CC in binding of leukocytes to the vascular endothelial wall prior to  
 CC extravasation of the leukocyte, e.g. to a site of inflammation.  
 CC The peptides bind pref. to E-selectin but may also bind L- or  
 CC P-selectin, and can be used to treat conditions mediated by E-selectin,  
 CC e.g. inflammatory conditions. The peptides have strong affinity for the  
 CC selectin receptors and inhibit the binding of the sialyl Lewis (SLe-x)  
 CC part of cell surface glycoproteins to E-selectin. The peptide are  
 CC small, generally less than 2 kD, have an IC50 of up to 100 micromole  
 CC against binding of HL60 cells to ELAM-1, have one or more peptide  
 CC linkages replaced by CH2OC(O)NR, phosphonate, CH2SO2NR, CON(R6),  
 CC or NHCONH linkages where R = H or a lower alkyl and R6 = a lower alkyl.

CC The peptides may also have substituted N- and C-termini e.g.  
 CC succinimido, N-benzoyloxycarbonyl or N-lower alkyl cpds.  
 XX  
 SQ Sequence 10 AA;  
 Query Match 68.9%; Score 31; DB 17; Length 10;  
 Best Local Similarity 100.0%; Pred. No. 56;  
 Matches 4; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 WVRW 4  
 | | | |  
 DB 6 WVRW 9

RESULT 17  
 AAR86145  
 ID AAR86145 standard; peptide; 10 AA.  
 XX AAR86145;  
 AC  
 XX  
 DT 26-JUN-1996 (first entry)  
 XX  
 DE Anti-ELAM-1 binding peptide #117.  
 XX  
 KW Peptide mimetic; endothelial leukocyte adhesion molecule; ELAM; selectin;  
 KW receptor; leukocyte; vascular wall; endothelium; extravasation;  
 KW inflammation; sialyl Lewis; cell surface glycoprotein; HL60 cell.  
 XX  
 OS Synthetic.  
 XX  
 FH Key Location/Qualifiers  
 FT Modified-site 10 /note= "contain amidated C-terminus"  
 FT  
 XX WO9531210-A1.  
 XX  
 PD 23-NOV-1995.  
 XX  
 PF 11-MAY-1995; 95WO-US06315.  
 XX  
 PR 11-MAY-1994; 94US-0241054.  
 XX  
 PA (AFFY-) AFFYMAX TECHNOLOGIES NV.  
 XX  
 PI Barrett RW, Cwirla SE, Dower WJ, Koller KJ, Lee J;  
 PI Martens CL, Ruhland-fritsch B;  
 XX  
 DR WPI; 1996-010687/01.  
 XX  
 PT New peptide(s) that bind to endothelial leukocyte adhesion molecule  
 PT 1 - useful for treating inflammation and other E-selectin mediated  
 PT diseases  
 XX  
 PS Disclosure; Page 17; 85pp; English.  
 XX  
 CC Peptides AAR86024-R86236 are examples of peptides and their mimetics  
 CC that bind to endothelial leukocyte adhesion molecule (ELAM)-1. This  
 CC molecule is a member of the selectin family of receptors and is involved  
 CC in binding of leukocytes to the vascular endothelial wall prior to  
 CC extravasation of the leukocyte, e.g. to a site of inflammation.  
 CC The peptides bind pref. to E-selectin but may also bind L- or  
 CC P-selectin, and can be used to treat conditions mediated by E-selectin,  
 CC e.g. inflammatory conditions. The peptides have strong affinity for the  
 CC selectin receptors and inhibit the binding of the sialyl Lewis (SLe-x)  
 CC part of cell surface glycoproteins to E-selectin. The peptide are  
 CC small, generally less than 2 kD, have an IC50 of up to 100 micromole  
 CC against binding of HL60 cells to ELAM-1, have one or more peptide  
 CC linkages replaced by CH2OC(O)NR, phosphonate, CH2SO2NR, CON(R6),  
 CC or NHCONH linkages where R = H or a lower alkyl and R6 = a lower alkyl.  
 CC The peptides may also have substituted N- and C-termini e.g.  
 CC succinimido, N-benzoyloxycarbonyl or N-lower alkyl cpds.  
 XX  
 SQ Sequence 10 AA;

Query Match 68.9%; Score 31; DB 17; Length 10;  
Best Local Similarity 100.0%; Pred. No. 56;  
Matches 4; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

OY 1 WVRW 4  
Db 6 WVRW 9

RESULT 18  
AAR86146  
ID AAR86146 standard; peptide; 10 AA.  
XX AC AAR86146;  
XX DT 26-JUN-1996 (first entry)  
XX DE Anti-ELAM-1 binding peptide #123.  
XX KW Peptide mimetic; endothelial leukocyte adhesion molecule; ELAM; selectin;  
KW receptor; leukocyte; vascular wall; endothelium; extravasation;  
KW inflammation; sialyl Lewis; cell surface glycoprotein; HL60 cell.  
XX OS Synthetic.  
XX PN WO9531210-A1.  
XX PD 23-NOV-1995.  
XX PF 11-MAY-1995; 95WO-US06315.  
XX PR 11-MAY-1994; 94US-0241054.  
XX PA (AFFY-) AFFYMAX TECHNOLOGIES NV.  
XX PI Barrett RW, Cwirla SE, Dower WJ, Koller KJ, Lee J;  
PI Martens CL, Ruhland-fritsch B;  
XX WPI; 1996-010687/01.  
XX New peptide(s) that bind to endothelial leukocyte adhesion molecule  
PT 1 - useful for treating inflammation and other E-selectin mediated  
PT diseases  
XX PS Disclosure; Page 17; 85pp; English.

Peptides AAR86024-R86236 are examples of peptides and their mimetics  
that bind to endothelial leukocyte adhesion molecule (ELAM)-1. This  
molecule is a member of the selectin family of receptors and is involved  
in binding of leukocytes to the vascular endothelial wall prior to  
extravasation of the leukocyte, e.g. to a site of inflammation.  
The peptides bind pref. to E-selectin but may also bind L- or  
P-selectin, and can be used to treat conditions mediated by E-selectin,  
e.g. inflammatory conditions. The peptides have strong affinity for the  
selectin receptors and inhibit the binding of the sialyl Lewis (SLe-x)  
part of cell surface glycoproteins to E-selectin. The peptide are  
small, generally less than 2 kD, have an IC50 of up to 100 micromole  
against binding of HL60 cells to ELAM-1, have one or more peptide  
linkages replaced by CH2OC(O)NR, phosphonate, CH2SO2NR, CON(R6),  
or NHCONH linkages where R = H or a lower alkyl and R6 = a lower alkyl.  
The peptides may also have substituted N- and C-terminal e.g.  
succinimido, N-benzoyloxycarbonyl or N-lower alkyl cpds.

XX Sequence 10 AA;

Query Match 68.9%; Score 31; DB 17; Length 10;  
Best Local Similarity 100.0%; Pred. No. 56;  
Matches 4; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

OY 1 WVRW 4  
Db 6 WVRW 9

RESULT 19  
AAW63963  
ID AAW63963 standard; peptide; 10 AA.  
XX AC AAW63963;  
XX DT 25-MAR-2003 (updated)  
DT 02-OCT-1998 (first entry)  
XX DE ELAM-1 peptide mimetic #118.  
XX KW Endothelial leukocyte adhesion molecule 1; ELAM-1; inflammation;  
KW selectin; diagnosis; mimetic.  
XX OS Synthetic.  
XX FH Key Location/Qualifiers  
FT Modified-site 10  
FT /note= "C-terminal Met is amidated"  
XX PN US5728802-A.  
XX PD 17-MAR-1998.  
XX PF 12-MAY-1995; 95US-0439817.  
XX PR 12-MAY-1995; 95US-0439817.  
PR 06-MAY-1992; 92US-0881395.  
PR 05-MAY-1993; 93US-0057295.  
PR 11-MAY-1994; 94US-0241054.  
XX PA (AFFY-) AFFYMAX TECHNOLOGIES NV.  
XX PI Barrett RW, Cwirla SE, Dower WJ, Koller KJ, Lee J;  
PI Martens CL, Ruhland B;  
XX WPI; 1998-249882/22.  
XX Peptide(s) or their mimetic(s) that bind to E-selectin - useful for,  
PT e.g. treating conditions mediated by E-selectin such as inflammatory  
PT condition(s)  
XX PS Example 2; Column 93-94; 84pp; English.

AAW63846-W64054 are peptides and peptide mimetics that bind selectins  
including endothelial leukocyte adhesion molecule 1 (ELAM-1) and can be  
used for blocking adhesion of leukocytes to the selectins. The peptides  
have applications for the treatment of conditions mediated by  
E-selectin, e.g. inflammatory conditions. They can also be used for  
diagnostic purposes, e.g. for identifying the vascular site of E-selectin  
in vivo or can be coupled to anti-inflammatory or other drugs.  
(Updated on 25-MAR-2003 to correct PF field.)

XX Sequence 10 AA;

Query Match 68.9%; Score 31; DB 19; Length 10;  
Best Local Similarity 100.0%; Pred. No. 56;  
Matches 4; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

OY 1 WVRW 4  
Db 6 WVRW 9

RESULT 20  
AAW63964  
ID AAW63964 standard; peptide; 10 AA.  
XX AC AAW63964;  
XX DT 25-MAR-2003 (updated)

DT 02-OCT-1998 (first entry)  
 XX ELAM-1 peptide mimetic #119.  
 DE Endothelial leukocyte adhesion molecule 1; ELAM-1; inflammation;  
 XX selectin; diagnosis; mimetic.  
 KW Synthetic.  
 KW US5728802-A.  
 XX PD 17-MAR-1998.  
 XX 12-MAY-1995; 95US-0439817.  
 XX 12-MAY-1995; 95US-0439817.  
 PR 06-MAY-1992; 92US-0881395.  
 PR 05-MAY-1993; 93US-0057295.  
 PR 11-MAY-1994; 94US-0241054.  
 XX (AFFY-) AFFYMAX TECHNOLOGIES NV.  
 PA Barrett RW, Cwirla SE, Dower WJ, Koller KJ, Lee J;  
 XX Martens CL, Ruhland B;  
 PI WPI; 1998-249882/22.  
 XX Peptide(s) or their mimetic(s) that bind to E-selectin - useful for,  
 PT e.g. treating conditions mediated by E-selectin such as inflammatory  
 PT condition(s)  
 PT Example 2; Column 93-94; 84pp; English.  
 PS AAW63846-W64054 are peptides and peptide mimetics that bind selectins  
 XX including endothelial leukocyte adhesion molecule 1 (ELAM-1) and can be  
 CC used for blocking adhesion of leukocytes to the selectins. The peptides  
 CC have applications for the treatment of conditions mediated by  
 CC E-selectin, e.g. inflammatory conditions. They can also be used for  
 CC diagnostic purposes, e.g. for identifying the vascular site of E-selectin  
 CC in vivo or can be coupled to anti-inflammatory or other drugs.  
 CC (Updated on 25-MAR-2003 to correct PF field.)  
 XX Query Match 68.9%; Score 31; DB 19; Length 10;  
 SQ Best Local Similarity 100.0%; Pred. No. 56;  
 Matches 4; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  
 QY 1 WVRW 4  
 DB |||||  
 6 WVRW 9  
 RESULT 21  
 AAW63958  
 ID AAW63958 standard; peptide; 10 AA.  
 AC AAW63958;  
 XX 25-MAR-2003 (updated)  
 DT 02-OCT-1998 (first entry)  
 XX ELAM-1 peptide mimetic #113.  
 DE Endothelial leukocyte adhesion molecule 1; ELAM-1; inflammation;  
 KW selectin; diagnosis; mimetic.  
 KW Synthetic.  
 OS US5728802-A.  
 XX PD 17-MAR-1998.  
 XX Query Match 68.9%; Score 31; DB 19; Length 10;  
 SQ Best Local Similarity 100.0%; Pred. No. 56;  
 Matches 4; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  
 QY 1 WVRW 4  
 DB |||||  
 6 WVRW 9  
 RESULT 22  
 AAR37390  
 ID AAR37390 standard; peptide; 6 AA.  
 AC AAR37390;  
 XX 07-JUL-1993 (first entry)  
 DT Peptide for treating septic shock.  
 DE Toxic shock; blood endotoxin removal; serum; diagnostic reagent;  
 XX cytokine release control; treatment; pertussis; bacterial meningitis;  
 KW HIV related infections; polymyxin B; Group II.  
 XX Synthetic.  
 OS Location/Qualifiers  
 XX Key 1..3  
 FT Region /note= "repeat region"  
 FT 4..6  
 FT Region /note= "repeat region"  
 FT ZAG200943-A.  
 PN 25-NOV-1992.  
 XX 10-FEB-1992; 92ZA-0000943.  
 PF 11-FEB-1991; 91US-0658744.  
 XX (PORR/) PORRO M.  
 PA Porro M;  
 PI

PF 12-MAY-1995; 95US-0439817.  
 XX 12-MAY-1995; 95US-0439817.  
 PR 06-MAY-1992; 92US-0881395.  
 PR 05-MAY-1993; 93US-0057295.  
 PR 11-MAY-1994; 94US-0241054.  
 XX (AFFY-) AFFYMAX TECHNOLOGIES NV.  
 PA Barrett RW, Cwirla SE, Dower WJ, Koller KJ, Lee J;  
 XX Martens CL, Ruhland B;  
 PI WPI; 1998-249882/22.  
 XX Peptide(s) or their mimetic(s) that bind to E-selectin - useful for,  
 PT e.g. treating conditions mediated by E-selectin such as inflammatory  
 PT condition(s)  
 PT Example 2; Column 91-92; 84pp; English.  
 PS AAW63846-W64054 are peptides and peptide mimetics that bind selectins  
 XX including endothelial leukocyte adhesion molecule 1 (ELAM-1) and can be  
 CC used for blocking adhesion of leukocytes to the selectins. The peptides  
 CC have applications for the treatment of conditions mediated by  
 CC E-selectin, e.g. inflammatory conditions. They can also be used for  
 CC diagnostic purposes, e.g. for identifying the vascular site of E-selectin  
 CC in vivo or can be coupled to anti-inflammatory or other drugs.  
 CC (Updated on 25-MAR-2003 to correct PF field.)  
 XX Query Match 68.9%; Score 31; DB 19; Length 10;  
 SQ Best Local Similarity 100.0%; Pred. No. 56;  
 Matches 4; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  
 QY 1 WVRW 4  
 DB |||||  
 6 WVRW 9  
 RESULT 22  
 AAR37390  
 ID AAR37390 standard; peptide; 6 AA.  
 AC AAR37390;  
 XX 07-JUL-1993 (first entry)  
 DT Peptide for treating septic shock.  
 DE Toxic shock; blood endotoxin removal; serum; diagnostic reagent;  
 XX cytokine release control; treatment; pertussis; bacterial meningitis;  
 KW HIV related infections; polymyxin B; Group II.  
 XX Synthetic.  
 OS Location/Qualifiers  
 XX Key 1..3  
 FT Region /note= "repeat region"  
 FT 4..6  
 FT Region /note= "repeat region"  
 FT ZAG200943-A.  
 PN 25-NOV-1992.  
 XX 10-FEB-1992; 92ZA-0000943.  
 PF 11-FEB-1991; 91US-0658744.  
 XX (PORR/) PORRO M.  
 PA Porro M;  
 PI

XX WPI; 1993-094304/11.

XX New peptide for treatment or prevention of toxic shock - comprises

PT specified sequences of aminoacid(s) and analogs

PT comprising sequences retro-orientated

XX Example; Page 5; 39pp; English.

XX The (Group II) peptide is an example of a generic peptide of formula

CC R-( Lys/Arg/His - Phe/Tyr/Trp - Leu/Ile/Val )n-R, where n = 1-100

CC and each R is H, an amino acid residue or a fatty acid residue.

CC The peptide is useful for treating or preventing septic shock,

CC mixing with polymyxin B to reduce its toxicity; removing

CC endotoxins from blood, sera or other fluids (in vivo or in

CC vitro); controlling release of cytokines induced by endotoxins;

CC as diagnostic reagents to detect and quantify toxins in blood

CC or sera; preparing non-toxic antigenic complexes of lipid A or

CC lipopolysaccharide (LPS); and for treating pertussis, bacterial

CC meningitis and HIV-related infections. The usual dose is 10-100

CC ug/kg/day, given parenterally. It binds to the same sites as

CC polymyxin B, i.e. it inhibits all the toxic effects of lipid A. It

CC has no antibiotic activity; does not lyse erythrocytes; has no

CC toxicity in mice when injected at 50mg/kg and is relatively unstable

CC against proteases.

XX

SQ Sequence 6 AA;

Query Match 66.7%; Score 30; DB 14; Length 6;

Best Local Similarity 75.0%; Pred. No. 9.3e+05;

Matches 3; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

QY 1 WVRW 4

Db |:||

2 WIRW 5

RESULT 23

AAW66066

ID AAW66066 standard; peptide; 6 AA.

XX

AC AAW66066;

XX

DT 16-NOV-1998 (first entry)

XX

DE Peptide useful as somatostatin antagonist.

XX

KW somatostatin antagonist; growth hormone; insulin; glucagon; diabetes;

KW growth promoter; gastric enzyme; eating disorder; disulphide.

XX

OS Synthetic.

XX

FH Key Location/Qualifiers

FT Misc-difference 1..6

FT /note= "D-form residues"

XX

PN EP863156-A1.

XX

PD 09-SEP-1998.

XX

PF 05-MAR-1998; 98EP-0301654.

XX

PR 06-MAR-1997; 97US-0812724.

XX

PA (AMCY ) AMERICAN CYANAMID CO.

XX

PI Baumbach WR, Houghten RA;

XX

DR WPI; 1998-458800/40.

XX

PT New somatostatin antagonist peptide(s) - useful as animal growth

PT promoters

XX

PS Example 3; Page 10; 37pp; English.

XX

CC The invention relates to somatostatin antagonists that can be used to

CC promote the growth of meat-producing animals by decreasing the effect of

CC somatostatin and/or increasing the release of growth hormone, insulin,

CC glucagon and/or gastric enzymes and/or enhancing immune function. Pure

CC somatostatin antagonists may also be useful for treating human or animal

CC disorders where reversal of somatostatin activity is beneficial, e.g.

CC gastrointestinal or eating disorders, diabetes or brain dysfunction. The

CC present sequence represents a somatostatin antagonist.

XX

SQ Sequence 6 AA;

Query Match 66.7%; Score 30; DB 19; Length 6;

Best Local Similarity 75.0%; Pred. No. 9.3e+05;

Matches 3; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

QY 1 WVRW 4

Db |:||

2 WIRW 5

RESULT 24

AAAY24292

ID AAAY24292 standard; peptide; 6 AA.

XX

AC AAAY24292;

XX

DT 15-SEP-1999 (first entry)

XX

DE Somatostatin antagonist peptide from US5925618 Example 3.

XX

KW Somatostatin antagonist; growth hormone; insulin; glucagon; gastric;

KW enzyme; immune function; cyclic peptide; gastrointestinal disorder;

KW eating disorder; diabetes; brain dysfunction.

XX

OS Synthetic.

XX

PN US5925618-A.

XX

PD 20-JUL-1999.

XX

PF 03-MAR-1998; 98US-0033395.

XX

PR 06-MAR-1997; 97US-0035181.

PR 03-MAR-1998; 98US-0033395.

XX

PA (AMCY ) AMERICAN CYANAMID CO.

XX

PI Baumbach WR, Houghten RA;

XX

DR WPI; 1999-429054/36.

XX

FT New peptides, used to treat gastrointestinal and eating disorders,

FT diabetes, and brain dysfunction

XX

PS Example 3; Column 10; 15pp; English.

XX

CC The present invention describes linear and cyclic peptides, which

CC decrease the effect of somatostatin. The somatostatin antagonist

CC peptides are used for decreasing the effect of somatostatin, by

CC contacting a somatostatin receptor site. They are also used for

CC increasing the release of insulin, increasing the release of glucagon,

CC enhancing the growth of animals and enhancing immune function. They can

CC be used to treat gastrointestinal and eating disorders, diabetes and

CC brain dysfunction, and also to increase growth in meat producing

CC animals. The peptides demonstrate inverse agonist activity. This allows

CC them to act as pure somatostatin antagonists, while blocking intrinsic

CC somatostatin receptor activity, independent of endogenous somatostatin.

CC AAAY24253 to AAAY24304 represent peptides used in the exemplification of

CC the present invention.

XX

SQ Sequence 6 AA;

Query Match 66.7%; Score 30; DB 20; Length 6;  
Best Local Similarity 75.0%; Pred. No. 9.3e+05;  
Matches 3; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

QY 1 WVRW 4  
|:|  
Db 2 WIRW 5

RESULT 25  
ABR45592  
ID ABR45592 standard; Peptide; 6 AA.  
XX AC ABR45592;  
XX DT 10-JUN-2003 (first entry)  
XX DE Staphylococcus aureus CHIPS-related peptide #782.  
XX KW CHIPS; Chemotaxis Inhibitory Protein; C5a-receptor; C5aR;  
KW formulated peptide receptor; FPR; neutrophil; monocyte; endothelial cell;  
KW inflammation; cardiovascular disease; central nervous system disease;  
KW gastrointestinal disease; skin disease; genitourinary disease;  
KW joint disease; respiratory disease; HIV infection; antiinflammatory;  
KW cardiant; cerebroprotective; neuroprotective; nootropic; dermatological;  
KW gynecological; immunosuppressive; anti-HIV.  
XX OS Staphylococcus aureus.  
OS Synthetic.  
XX WO2003006048-A1.  
XX PD 23-JAN-2003.  
XX PF 11-JUL-2001; 2001WO-EP08004.  
XX PR 11-JUL-2001; 2001WO-EP08004.  
XX PA (JARI-) JARI PHARM BV.  
XX PI Van Kessel CPM, Gosselaar-de Haas CJC, Kruijtzer JAW;  
PI Van Strijp JAG;  
XX DR WPI; 2003-247783/25.  
XX PT Combination of peptides derived from chemotaxis inhibiting protein from  
PT Staphylococcus aureus (CHIPS) having CHIPS activity, useful in  
PT prophylaxis and treatment of inflammation, cardiovascular, skin and  
PT kidney diseases  
XX PS Disclosure; Page 13; 89pp; English.  
XX CC The present invention relates to peptides (ABR44811-ABR47162 and  
CC ABR47164-ABR47385) derived from the Chemotaxis Inhibitory Protein (CHIPS)  
CC from Staphylococcus aureus. The peptide fragments are useful in the  
CC prophylaxis or treatment of diseases or disorders involving the  
CC C5a-receptor (C5aR) and/or formulated peptide receptor (FPR) or  
CC neutrophils, monocytes and endothelial cells or involving acute or  
CC chronic inflammation reactions. The diseases or disorders include  
CC cardiovascular diseases, disease of the central nervous system,  
CC gastrointestinal diseases, skin diseases, genitourinary diseases, joint  
CC diseases, respiratory diseases and HIV infection.  
XX SQ Sequence 6 AA;  
Query Match 66.7%; Score 30; DB 24; Length 6;  
Best Local Similarity 50.0%; Pred. No. 9.3e+05;  
Matches 3; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

QY 1 WVRWHF 6  
|:|  
Db 1 WIFWYF 6

RESULT 26  
AAY08189  
ID AAY08189 standard; peptide; 8 AA.  
XX AC AAY08189;  
XX DT 09-JUL-1999 (first entry)  
XX DE Clotting factor VIII binding peptide 71.  
XX KW Coagulation factor VIII; clotting factor VIII; diagnosis; treatment;  
KW purification; disorder; blood coagulation.  
XX OS Synthetic.  
XX PN WO9914232-A1.  
XX PD 25-MAR-1999.  
XX PF 12-SEP-1998; 98WO-EP05822.  
XX PR 13-SEP-1997; 97DE-1040310.  
XX PA (OCTA-) OCTAPHARMA AG.  
XX PI Jungbauer A;  
XX DR WPI; 1999-312410/26.  
XX PT Peptides with affinity for blood clotting factor 8  
XX PS Claim 4; Page 38; 51pp; German.  
XX CC This invention describes novel peptides (AAY08119-Y08212) with affinity  
CC for coagulation factor VIII which can be used for for labeling,  
CC identification (diagnostic) and purification of factor VIII. Some are  
CC specific for one of natural and recombinant factor VIII, others are  
CC reactive with both forms. Factor VIII is used to treat disorders of  
CC blood coagulation. Using relatively small peptides, rather than large  
CC antibody molecules generally used, simplifies purification of factor  
CC VIII. The peptides are of formula R1-X-R2 where R1 = amino or a  
CC peptide; R2 = carboxy or a peptide and X = a peptide of at least 3,  
CC preferably 7-12, amino acid residues.  
XX SQ Sequence 8 AA;  
Query Match 66.7%; Score 30; DB 20; Length 8;  
Best Local Similarity 33.3%; Pred. No. 9.3e+05;  
Matches 2; Conservative 3; Mismatches 1; Indels 0; Gaps 0;

QY 1 WVRWHF 6  
|:|  
Db 2 WIKWEY 7

RESULT 27  
AAW80380  
ID AAW80380 standard; Peptide; 12 AA.  
XX AC AAW80380;  
XX DT 14-JAN-1999 (first entry)  
XX DE Peptide eluted after biopanning against maltose binding protein.  
XX KW Intervening protein sequence; IVPS; protein splicing;  
KW protein production; maltose binding protein.  
XX OS Synthetic.  
XX PN US5834247-A.



XX 10-NOV-1998.  
 XX PD  
 XX PF  
 XX PI  
 XX XX  
 PR 05-MAR-1997; 97US-0811492.  
 PR 05-MAR-1997; 97US-0811492.  
 PR 09-DEC-1992; 92US-0004139.  
 PR 03-NOV-1993; 93US-0146885.  
 PR 28-JUN-1995; 95US-0496247.  
 PR 29-DEC-1995; 95US-0580555.  
 XX XX  
 PA (NEW ) NEW ENGLAND BIOLABS INC.  
 XX Adam E, Chong SSC, Comb DG, Hodges RA, Jack WE;  
 PI Noren CJ, Perler FB, Southworth M, Xu M;  
 XX WPI; 1999-008713/01.  
 XX New modified target proteins - which have controllable intervening  
 PT protein sequence which can facilitate production, purification,  
 PT labelling or isolation of target proteins  
 XX  
 PS Example 22; Fig 36; 123pp; English.  
 XX  
 CC AAW80372-93 represent peptides eluted after biopanning against  
 CC maltose binding protein, in the course of the invention. The  
 CC specification describes IVPS (intervening protein sequence)  
 CC regions which encode peptides which are removed via protein  
 CC splicing to form the native protein. The specification describes  
 CC a modified protein comprising a target protein or portion, fused  
 CC either internally or terminally, to a IVPS, or to an amino- or  
 CC carboxyl-terminal element of a IVPS. The IVPS are capable of  
 CC excision from or cleavage of the modified protein upon predetermined  
 CC conditions, in cis or trans, e.g. temperature increase, deglycosylation,  
 CC unblocking of amino acid residues, treatment with chemical reagents.  
 CC The methods can be used for modifying, producing, purifying, labelling  
 CC or isolating target proteins such as enzymes, toxins, cytokines,  
 CC glycoproteins and growth factors.  
 XX  
 SQ Sequence 12 AA;  
 Query Match 66.7%; Score 30; DB 20; Length 12;  
 Best Local Similarity 100.0%; Pred. No. 97;  
 Matches 4; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  
 QY 3 RWHF 6  
 Db ||||  
 7 RWHF 10  
 RESULT 28  
 ABB74383  
 ID ABB74383 standard; Peptide; 14 AA.  
 XX  
 AC ABB74383;  
 XX  
 DT 18-APR-2002 (first entry)  
 XX  
 DE Karyophilic peptide SEQ ID NO:147.  
 XX  
 KW Fusogenic; nuclear localisation signal; NLS; encapsulation; lipogene;  
 KW liposome; micelle; karyophilic; cytostatic; antitumour; solid tumour;  
 KW peptide-lipid-polynucleotide complex; neoplastic disease; gene therapy;  
 KW breast carcinoma; prostate carcinoma.  
 XX  
 OS Saccharomyces cerevisiae.  
 XX  
 PN WO200193836-A2.  
 XX  
 PD 13-DEC-2001.  
 XX  
 PF 08-JUN-2001; 2001WO-US18657.  
 XX

PR 09-JUN-2000; 2000US-210925P.  
 XX (BOUL/) BOULIKAS T.  
 XX  
 XX PI  
 XX Boulukas T;  
 DR WPI; 2002-164295/21.  
 XX  
 PT Encapsulation of plasmid DNA (Lipogenes) and therapeutic agents with  
 PT nuclear localization signal/fusogenic peptide conjugates into targeted  
 PT liposome complexes -  
 XX  
 PS Claim 14; Page 63; 107pp; English.  
 XX  
 CC The present invention describes a method for producing micelles with  
 CC entrapped therapeutic agents. The method comprises: (1) combining  
 CC negatively charged agent with a cationic lipid in a ratio where 30-90 %  
 CC of the negatively charged atoms are neutralised by positive charges on  
 CC lipid molecules to form an electrostatic micelle complex in 20-80 %  
 CC ethanol; and (2) combining the micelle complex of (a) with fusogenic-  
 CC karyophilic peptide conjugates in a 0.0-0.3 ratio, therefore producing  
 CC micelles with entrapped therapeutic agents. Also described is a method  
 CC for delivering a therapeutic agent in vivo, comprising the administration  
 CC of the micelle. ABB74256 to ABB74858 represent specifically claimed  
 CC nuclear localisation signal (NLS) peptides for use in the method as the  
 CC fusogenic-karyophilic peptides. The micelles produced can have cytostatic  
 CC and antitumour activities. The peptide-lipid-polynucleotide complexes  
 CC produced are useful for inhibiting the progression of neoplastic  
 CC diseases. The invention relates to the field of gene therapy and is  
 CC directed toward methods for producing peptide-lipid-polynucleotide  
 CC complexes suitable for delivery of polynucleotides. The encapsulated  
 CC molecules display therapeutic efficacy in eradicating solid tumours  
 CC including but not limited to breast carcinoma or prostate carcinoma.  
 CC ABB74235 to ABB74255 are used in the exemplification of the present  
 CC invention.  
 XX  
 SQ Sequence 14 AA;  
 Query Match 66.7%; Score 30; DB 23; Length 14;  
 Best Local Similarity 100.0%; Pred. No. 1.1e+02;  
 Matches 4; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  
 QY 3 RWHF 6  
 Db ||||  
 10 RWHF 13  
 RESULT 29  
 AAB49729  
 ID AAB49729 standard; peptide; 7 AA.  
 XX  
 AC AAB49729;  
 XX  
 DT 10-APR-2001 (first entry)  
 XX  
 DE Peptide SEQ ID 40 which binds to the TADG5 protein.  
 XX  
 KW TADG5; human; zinc finger; SH3 domain; cell signalling;  
 KW cell cycle control.  
 XX  
 OS Unidentified.  
 XX  
 PN WO200102432-A1.  
 XX  
 PD 11-JAN-2001.  
 XX  
 PF 30-JUN-2000; 2000WO-US18304.  
 XX  
 PR 01-JUL-1999; 99US-0346510.  
 XX  
 PA (UYAR-) UNIV ARKANSAS.  
 XX  
 PI O'Brien TJ, Wang Y;

XX WPI; 2001-123102/13.

DR Novel SH3 domain-containing TADG5 protein useful for regulating gene

XX replication, as a nutrition supplement, and as a marker for human

PT tissue, or in cell cycle control -

PT Example 6; Page 36; 85pp; English.

XX This invention relates to an SH3 domain-containing protein termed TADG5,

CC and its variants. The invention includes amino acid and polynucleotide

CC sequences for TADG5, and oligonucleotides which bind to either the basic

CC amino acid region and/or the zinc finger motif of the TADG5 protein. The

CC basic amino acid region or zinc finger motif of TADG5 is useful for

CC regulating the expression of the TADG5 gene in a cell. The TADG5 protein

CC is useful as a source of amino acids, as a nutrition supplement, and as a

CC marker for human tissue, or in cell cycle control. TADG5 protein or

CC peptides generated from the protein sequence are useful as antigens for

CC the production of polyclonal and monoclonal antibodies. DNA encoding

CC TADG5 is useful as an antisense vehicle for cell cycle control by

CC shutting down signalling or cell division. The present sequence

CC represents a peptide identified from a phage display peptide library

CC through biopanning with the TADG5 protein.

XX

SQ Sequence 7 AA;

Query Match 64.4%; Score 29; DB 22; Length 7;

Best Local Similarity 60.0%; Pred. No. 9.3e+05;

Matches 3; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

OY 1 WVRWH 5

Db 3 WMDWH 7

RESULT 30

ABB90493

ID ABB90493 standard; Peptide; 8 AA.

XX

AC ABB90493;

XX

DT 27-MAY-2002 (first entry)

XX

DE Hominidae LDL receptor related peptide sequence #139.

XX

KW Hominidae; low density lipoprotein receptor; LDL receptor; LDL-R;

KW detection; lipid metabolic error; hyperlipaemia; mutation;

KW arteriosclerosis; ischaemic heart disease; ischaemia.

XX

OS Hominidae.

OS Synthetic.

XX

PN WO200206467-A1.

XX

PD 24-JAN-2002.

XX

PF 17-JUL-2001; 2001WO-JP06153.

XX

PR 18-JUL-2000; 2000JP-0218039.

XX

PA (BMLB-) BML INC.

XX

PI Hattori H, Tsuji M, Okada T, Nagano M, Egashira T, Ishihara M;

PI Iwasaki T;

XX

DR WPI; 2002-179794/23.

XX

PT Set of specific low density lipoprotein receptor gene mutations for

PT diagnosis of familial lipid metabolism errors including hyperlipemia -

XX

PS Example; Fig 50; 123pp; Japanese.

XX

CC The present invention describes a method for detecting lipid metabolism

CC errors in patients using as indicators a set of 65 specific low density

CC lipoprotein (LDL) receptor gene mutations. The method can be used in the

CC diagnosis of an inherited predisposition to the development of diseases

CC associated with hyperlipaemia, such as arteriosclerosis and ischaemic

CC heart disease. ABL91141 encodes the LDL receptor given in ABB90525.

CC ABL91142 to ABL91183 represent PCR primers used in the amplification of

CC the receptor gene. ABL90990 to ABL91140 and ABB90445 to ABB90524

CC represents sequences used in the exemplification of the present

CC invention.

XX

SQ Sequence 8 AA;

Query Match 64.4%; Score 29; DB 23; Length 8;

Best Local Similarity 60.0%; Pred. No. 9.3e+05;

Matches 3; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

OY 1 WVRWH 5

Db 2 WPQWH 6

RESULT 31

AAR33522

ID AAR33522 standard; peptide; 6 AA.

XX

AC AAR33522;

XX

DT 07-JUL-1993 (first entry)

XX

DE Peptide for treating septic shock.

XX

KW Toxic shock; blood endotoxin removal; serum; diagnostic reagent;

KW cytokine release control; treatment; pertussis; bacterial meningitis;

KW HIV related infections; polymyxin B; Group I.

XX

OS Synthetic.

XX

FH Key Location/Qualifiers

FT Region 1..3

FT /note= "repeat region"

FT Region 4..6

FT /note= "repeat region"

XX

PN ZA9200943-A.

XX

PD 25-NOV-1992.

XX

PF 10-FEB-1992; 92ZA-0000943.

XX

PR 11-FEB-1991; 91US-0658744.

XX

PA (PORR/) PORRO M.

XX

PI Porro M;

XX

DR WPI; 1993-094304/11.

XX

PT New peptide for treatment or prevention of toxic shock - comprises

PT specified sequences of aminoacid(s) and analogs

PT comprising sequences retro-orientated

XX

PS Example; Page 5; 39pp; English.

XX

CC The (Group I) peptide is an example of a generic peptide of formula

CC R-( Lys/Arg/His - Phe/Tyr/Trp - Leu/Ile/Val )n-R, where n = 1-100

CC and each R is H, an amino acid residue or a fatty acid residue.

CC The peptide is useful for treating or preventing septic shock,

CC mixing with polymyxin B to reduce its toxicity; removing

CC endotoxins from blood, sera or other fluids (in vivo or in

CC vitro); controlling release of cytokines induced by endotoxins;

CC as diagnostic reagents to detect and quantify toxins in blood

CC or sera; preparing non-toxic antigenic complexes of lipid A or

CC lipopolysaccharide (LPS); and for treating pertussis, bacterial

CC meningitis and HIV-related infections. The usual dose is 10-100  
CC ug/kg/day, given parenterally. It binds to the same sites as  
CC polymyxin B, i.e. it inhibits all the toxic effects of lipid A. It  
CC has no antibiotic activity; does not lyse erythrocytes; has no  
CC toxicity in mice when injected at 50mg/kg and is relatively unstable  
CC against proteases.

XX Sequence 6 AA;  
SQ Query Match 62.2%; Score 28; DB 14; Length 6;  
Best Local Similarity 75.0%; Pred. No. 9.3e+05;  
Matches 3; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

QY 1 WVRW 4  
Db 2 WVKW 5

RESULT 32  
AAR37388  
ID AAR37388 standard; peptide; 6 AA.  
XX  
AC AAR37388;  
XX  
DT 07-JUL-1993 (first entry)  
XX  
DE Peptide for treating septic shock.  
XX  
KW Toxic shock; blood endotoxin removal; serum; diagnostic reagent;  
KW cytokine release control; treatment; pertussis; bacterial meningitis;  
KW HIV related infections; polymyxin B; Group II.  
XX  
OS Synthetic.  
XX  
FH Key Location/Qualifiers  
FT Region 1..3 /note= "repeat region"  
FT Region 4..6 /note= "repeat region"  
FT  
FT  
FT  
XX  
PN ZA9200943-A.  
XX  
PD 25-NOV-1992.  
XX  
PF 10-FEB-1992; 92ZA-0000943.  
XX  
PR 11-FEB-1991; 91US-0658744.  
XX  
PA (PORR/) PORRO M.  
XX  
PI Porro M;  
XX  
DR WPI; 1993-094304/11.  
XX  
PT New peptide for treatment or prevention of toxic shock - comprises  
PT specified sequences of aminoacid(s) and analogs  
PT comprising sequences retro-orientated  
XX  
PS Example; Page 5; 39pp; English.  
XX  
CC The (Group II) peptide is an example of a generic peptide of formula  
CC R-( Lys/Arg/His - Phe/Tyr/Trp - Leu/Ile/Val )n-R, where n = 1-100  
CC and each R is H, an amino acid residue or a fatty acid residue.  
CC The peptide is useful for treating or preventing septic shock,  
CC mixing with polymyxin B to reduce its toxicity; removing  
CC endotoxins from blood, sera or other fluids (in vivo or in  
CC vitro); controlling release of cytokines induced by endotoxins;  
CC as diagnostic reagents to detect and quantify toxins in blood  
CC or sera; preparing non-toxic antigenic complexes of lipid A or  
CC lipopolysaccharide (LPS); and for treating pertussis, bacterial  
CC meningitis and HIV-related infections. The usual dose is 10-100  
CC ug/kg/day, given parenterally. It binds to the same sites as  
CC polymyxin B, i.e. it inhibits all the toxic effects of lipid A. It

CC has no antibiotic activity; does not lyse erythrocytes; has no  
CC toxicity in mice when injected at 50mg/kg and is relatively unstable  
CC against proteases.

XX Sequence 6 AA;  
SQ Query Match 62.2%; Score 28; DB 14; Length 6;  
Best Local Similarity 75.0%; Pred. No. 9.3e+05;  
Matches 3; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

QY 1 WVRW 4  
Db 2 WLRW 5

RESULT 33  
AAR93719  
ID AAR93719 standard; peptide; 6 AA.  
XX  
AC AAR93719;  
XX  
DT 10-MAY-1996 (first entry)  
XX  
DE Cyclo[-Tyr-trp-Leu-Arg-Trp-Pro-].  
XX  
KW neurokinin A antagonist; tachykinin; respiratory disease; asthma;  
KW analgesic; cyclic.  
XX  
OS Synthetic.  
XX  
FH Key Location/Qualifiers  
FT Modified-site 1 /note= "not an N-terminal amino acid, but condensed  
FT with Pro(6) to form a cyclic peptide"  
FT Misc-difference 2 /note= "D-form residue"  
FT Modified-site 6 /note= "not a C-terminal amino acid, but condensed  
FT with Tyr(1) to form a cyclic peptide"  
FT  
FT  
XX  
PN W09521187-A1.  
XX  
PD 10-AUG-1995.  
XX  
PF 10-JAN-1995; 95WO-US00296.  
XX  
PR 03-FEB-1994; 94US-0191571.  
XX  
PA (RICH ) MERRELL DOW PHARM INC.  
XX  
PI Buck SH, Harbeson SL, Kudlacz EM, Owen TJ;  
XX  
DR WPI; 1995-336695/43.  
XX  
PT New cyclic peptide derivs. - are neurokinin A and tachykinin  
PT antagonists useful e.g. for treating asthma or as analgesics  
XX  
PS Claims 25, 33; Pages 74, 76; 82pp; English.  
XX  
CC The patent describes novel cyclic hexapeptide and octapeptide compounds  
CC which are antagonists of neurokinin A and which are useful medically as  
CC analgesics and for treating respiratory diseases such as asthma. The  
CC patent also discloses the new use of a broader range of cyclic  
CC hexapeptides as analgesics and for treating respiratory diseases such  
CC as asthma. The present sequence represents a specifically preferred  
CC example of the broader peptides.  
XX  
SQ Sequence 6 AA;  
Query Match 62.2%; Score 28; DB 16; Length 6;  
Best Local Similarity 75.0%; Pred. No. 9.3e+05;  
Matches 3; Conservative 1; Mismatches 0; Indels 0; Gaps 0;



FT Modified-site 6 with Ala(6) to form a cyclic peptide"  
FT /note= "not a C-terminal amino acid, but condensed  
FT with Tyr(1) to form a cyclic peptide"  
FT Misc-difference 2  
FT /note= "D-form residue"  
FT Misc-difference 6  
FT /note= "L- or D-form residue"

XX WO9521187-A1.  
PN 10-AUG-1995.  
XX 10-JAN-1995; 95WO-US00296.  
PF 03-FEB-1994; 94US-0191571.  
XX (RICH ) MERRELL DOW PHARM INC.  
PR Buck SH, Harbeson SL, Kudlacz EM, Owen TJ;  
XX WPI; 1995-336695/43.

XX New cyclic peptide derivs. - are neurokinin A and tachykinin  
PT antagonists useful e.g. for treating asthma or as analgesics  
PT Claim 6; Page 69; 82pp; English.

XX The patent describes novel cyclic hexapeptide and octapeptide compounds  
CC which are antagonists of neurokinin A and which are useful medically as  
CC analgesics and for treating respiratory diseases such as asthma. The  
CC present sequence represents a specifically preferred example of the new  
CC peptides.

XX Sequence 6 AA;

Query Match 62.2%; Score 28; DB 16; Length 6;  
Best Local Similarity 75.0%; Pred. No. 9.3e+05;  
Matches 3; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

QY 1 WVRW 4  
Db 2 WLRW 5

RESULT 37  
AAR74033  
ID AAR74033 standard; Peptide; 10 AA.  
XX AAR74033;  
AC 19-DEC-1995 (first entry)  
XX Bombesin-related peptide SAP bombesin-10.  
DE Bombesin; frog; PCR; primer; amplification; probe; prohormone; human;  
XX veterinary medicine.  
KW Synthetic.

OS Key Location/Qualifiers  
XX Misc-difference 10  
FT /note= "amidated C-terminus"  
FT US5410018-A.  
XX 25-APR-1995.  
PD 25-FEB-1994; 94US-0203196.  
XX 25-FEB-1994; 94US-0203196.  
PF (OREG-) OREGON REGIONAL PRIMATE RES CENT.

XX Barry B, Nagalla S, Spindel ER;  
PI WPI; 1995-169632/22.  
XX Purified bombesin-related peptide(s) - prepared by recombinant DNA  
PT methods  
XX Claim 2; Column 7-8; 10pp; English.  
XX The peptides AAR74032-3 are derived from the bombesin-related prohormone  
CC AAR74034. The peptides are generated by internal processing of the  
CC prohormone at the Ser-Leu and Lys-Lys sequences. This peptide is  
CC designated SAP bombesin-10 ("BIM-26336") and corresponds to residues  
CC 49-58 of the prohormone. The SAP bombesin-10 is then modified from the  
CC prohormone-cleaved peptide by having an amidated methionyl residue.  
CC This peptide can be generated by an internal cleavage of the SAP  
CC bombesin-14 (AAR74032). The amide gp. being donated from the Gly residue  
CC at position 59 of the prohormone. The peptides have applications within  
CC human and veterinary medicine, especially to treat the diseases or  
CC disorders specified in US5217955, WO9402018 and WO9220363.

XX Sequence 10 AA;

Query Match 62.2%; Score 28; DB 16; Length 10;  
Best Local Similarity 66.7%; Pred. No. 1.7e+02;  
Matches 4; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

QY 1 WVRWHF 6  
Db 4 WARGHF 9

RESULT 38  
AAR86144  
ID AAR86144 standard; peptide; 10 AA.

XX AAR86144;  
AC 26-JUN-1996 (first entry)  
XX Anti-ELAM-1 binding peptide #121.

DE Peptide mimetic; endothelial leukocyte adhesion molecule; ELAM; selectin;  
XX receptor; leukocyte; vascular wall; endothelium; extravasation;  
KW inflammation; sialyl Lewis; cell surface glycoprotein; HL60 cell.

XX Synthetic.

XX WO9531210-A1.

XX 23-NOV-1995.

XX 11-MAY-1995; 95WO-US06315.

XX 11-MAY-1994; 94US-0241054.

XX (AFFY-) AFFYMAX TECHNOLOGIES NV.

PI Barrett RW, Cwirila SE, Dower WJ, Koller KJ, Lee J;  
PI Martens CL, Ruhland-fritsch B;

XX WPI; 1996-010687/01.

XX New peptide(s) that bind to endothelial leukocyte adhesion molecule  
PT 1 - useful for treating inflammation and other E-selectin mediated  
PT diseases

XX Disclosure; Page 17; 85pp; English.

XX Peptides AAR86024-R86236 are examples of peptides and their mimetics  
CC that bind to endothelial leukocyte adhesion molecule (ELAM)-1. This  
CC molecule is a member of the selectin family of receptors and is involved



CC in binding of leukocytes to the vascular endothelial wall prior to  
CC extravasation of the leukocyte, e.g. to a site of inflammation.  
CC The peptides bind pref. to E-selectin but may also bind L- or  
CC P-selectin, and can be used to treat conditions mediated by E-selectin,  
CC e.g. inflammatory conditions. The peptides have strong affinity for the  
CC selectin receptors and inhibit the binding of the sialyl Lewis (SLe-x)  
CC part of cell surface glycoproteins to E-selectin. The peptide are  
CC small, generally less than 2 kD, have an IC50 of up to 100 micromole  
CC against binding of HL60 cells to ELAM-1, have one or more peptide  
CC linkages, replaced by CH2OC(O)NR, phosphonate, CH2SO2NR, CH2NR, CON(R6),  
CC or NHCONH linkages where R = H or a lower alkyl and R6 = a lower alkyl.  
CC The peptides may also have substituted N- and C-termini e.g.  
CC succinimido, N-benzylloxycarbonyl or N-lower alkyl cpds.

XX SQ Sequence 10 AA;

Query Match 62.2%; Score 28; DB 17; Length 10;  
Best Local Similarity 75.0%; Pred. No. 1.7e+02;  
Matches 3; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

Oy 1 WVRW 4  
||:|  
Db 6 WVKW 9

RESULT 39

AAW63962  
ID AAW63962 standard; peptide; 10 AA.

XX AC AAW63962;

XX DT 25-MAR-2003 (updated)

DT 02-OCT-1998 (first entry)

XX DE ELAM-1 peptide mimetic #117.

XX KW Endothelial leukocyte adhesion molecule 1; ELAM-1; inflammation;  
KW selectin; diagnosis; mimetic.

XX OS Synthetic.

PN US5728802-A.

XX PD 17-MAR-1998.

XX PF 12-MAY-1995; 95US-0439817.

XX PR 12-MAY-1995; 95US-0439817.

PR 06-MAY-1992; 92US-0881395.

PR 05-MAY-1993; 93US-0057295.

PR 11-MAY-1994; 94US-0241054.

XX PA (AFFY-) AFFYMAX TECHNOLOGIES NV.

XX PI Barrett RW, Cwirla SE, Dower WJ, Koller KJ, Lee J;

PI Martens CL, Ruhland B;

XX DR WPI; 1998-249882/22.

XX PT Peptide(s) or their mimetic(s) that bind to E-selectin - useful for,  
PT e.g. treating conditions mediated by E-selectin such as inflammatory  
PT condition(s)

XX PS Example 2; Column 91-92; 84pp; English.

XX CC AAW63846-W64054 are peptides and peptide mimetics that bind selectins  
CC including endothelial leukocyte adhesion molecule 1 (ELAM-1) and can be  
CC used for blocking adhesion of leukocytes to the selectins. The peptides  
CC have applications for the treatment of conditions mediated by  
CC E-selectin, e.g. inflammatory conditions. They can also be used for  
CC diagnostic purposes, e.g. for identifying the vascular site of E-selectin  
CC in vivo or can be coupled to anti-inflammatory or other drugs.  
CC (Updated on 25-MAR-2003 to correct PF field.)

XX SQ Sequence 10 AA;  
Query Match 62.2%; Score 28; DB 19; Length 10;  
Best Local Similarity 75.0%; Pred. No. 1.7e+02;  
Matches 3; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

Oy 1 WVRW 4  
||:|  
Db 6 WVKW 9

RESULT 40

AAR36519  
ID AAR36519 standard; peptide; 12 AA.

XX AC AAR36519;

XX DT 25-MAR-2003 (updated)

DT 11-AUG-1993 (first entry)

XX DE D32.39 antibody isolated peptide.

XX KW Generation; screening; selection; screening; peptide ligands;  
KW receptor molecules; therapeutics; diagnostics.

XX OS Synthetic.

XX PH Key Location/Qualifiers

FT Region 3..8

FT /note= "D32.39 epitope"

XX PN WO9308278-A1.

XX PD 29-APR-1993.

XX PF 15-OCT-1992; 92WO-US08879.

XX PR 16-OCT-1991; 91US-0778233.

XX PA (AFFY-) AFFYMAX TECHNOLOGIES NV.

XX PI Cull MG, Miller JF, Schatz PJ, Stemmer WPC;

XX DR WPI; 1993-152471/18.

XX PT Random peptide library and screening method - using vectors  
PT encoding fusion proteins of DNA binding protein and peptide, used  
PT in screening for ligands

XX PS Disclosure; Fig 3; 153pp; English.

XX CC The sequence is that of a peptide isolated by panning with the  
CC D32.39 antibody, with an ELISA result of 0.1. This was done as  
CC an example of a method of constructing a random peptide library  
CC of at least 10(6) members. The method enables the generation,  
CC screening and selection of peptide ligands and receptor molecules.  
CC Peptides generated using the method can be used in therapeutics  
CC and diagnostics, e.g. to inhibit receptor activity.  
CC (Updated on 25-MAR-2003 to correct PN field.)

XX SQ Sequence 12 AA;

Query Match 62.2%; Score 28; DB 14; Length 12;  
Best Local Similarity 100.0%; Pred. No. 2e+02;  
Matches 4; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Oy 2 VRWH 5

Db 2 VRWH 5

RESULT 41

AAR56756  
ID AAR56756 standard; peptide; 12 AA.  
AC AAR56756;  
XX  
DT 25-MAR-2003 (updated)  
DT 20-MAR-1995 (first entry)  
XX  
DE Random peptide #53 isolated by anti-dynorphin B Ab panning.  
XX  
KW Dynorphin B; epitope; antibody panning; random peptide library;  
KW antibody D32.39; ligand screening.  
XX  
OS Synthetic.  
XX  
FH Key Location/Qualifiers  
FT Peptide 3..8  
FT /note= "D32.39 epitope"  
XX  
PN US5498530-A.  
XX  
PD 12-MAR-1996.  
XX  
PF 15-AUG-1994; 94US-0290641.  
XX  
PR 15-OCT-1992; 92US-0963321.  
PR 16-OCT-1991; 91US-0778233.  
PR 15-AUG-1994; 94US-0290641.  
XX  
PA (AFFY-) AFFYMAX TECHNOLOGIES NV.  
XX  
PI Cull MG, Miller JF, Schatz PJ, Stemmer WPC;  
XX  
DR WPI; 1996-159686/16.  
XX  
PT Random peptide libraries comprising host cells expressing DNA  
PT binding proteins fused with random peptide(s) - used to identify,  
PT e.g. peptide ligands of receptors  
XX  
PS Example 4; Fig 3B; 45pp; English.  
XX  
CC Construction of random peptide library - by creating vectors  
CC contg. DNA encoding the random peptide(s) fused to DNA binding  
CC proteins; used to screen for novel ligands  
XX  
PS Example 4; Fig 3B; 45pp; English.  
XX  
CC A random peptide library was constructed in E.coli hosts.  
CC The library was lysed and panned using antibody D32.39 which  
CC recognises the Dynorphin B epitope RQFKV. Peptides isolated by  
CC panning were sequenced and a consensus epitope was identified (see  
CC features table). Arginine is invariant in the first position for all  
CC the ELISA positive clones (AAR56701-R56758). No strong bias was  
CC evident for the second position but in the third position, 5 amino  
CC acids (Phe, His, Asp, Tyr, Trp) account for 98% of the residues. The  
CC fourth position shows a strong bias for positively charged residues  
CC (Lys and Arg) with almost exclusively hydrophobic residues at  
CC position 5 (mostly Val). Val and Thr predominate at the sixth  
CC position (76%) with Ser and Ile accounting for the remaining amino  
CC acids.  
CC (Updated on 25-MAR-2003 to correct PF field.)  
XX  
SQ Sequence 12 AA;  
Query Match 62.2%; Score 28; DB 15; Length 12;  
Best Local Similarity 100.0%; Pred. No. 2e+02;  
Matches 4; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  
QY 2 VRWH 5  
DB |||||  
2 VRWH 5  
RESULT 42  
AAR91504  
ID AAR91504 standard; Peptide; 12 AA.  
XX  
AC AAR91504;  
XX  
DT 25-MAR-2003 (updated)

21-NOV-1996 (first entry)  
D32.39 monoclonal antibody peptide ligand 53.  
dynorphin B; random peptide library; construction; monoclonal antibody;  
D32.39; epitope; screening.  
Synthetic.  
Key Location/Qualifiers  
Peptide 3..8  
/note= "D32.39 epitope"  
US5498530-A.  
12-MAR-1996.  
15-AUG-1994; 94US-0290641.  
15-OCT-1992; 92US-0963321.  
16-OCT-1991; 91US-0778233.  
15-AUG-1994; 94US-0290641.  
(AFFY-) AFFYMAX TECHNOLOGIES NV.  
Cull MG, Miller JF, Schatz PJ, Stemmer WPC;  
WPI; 1996-159686/16.  
Random peptide libraries comprising host cells expressing DNA  
binding proteins fused with random peptide(s) - used to identify,  
e.g. peptide ligands of receptors  
Example 4; Fig 3B; 46pp; English.  
A random peptide (RP) library can be constructed by transforming host  
cells with a collection of recombinant vectors that encode a fusion  
protein comprised of a DNA binding protein (BP) and a RP and also  
contains a binding site for the DNA BP. The RP library can be used to  
screen for novel ligands, the method resulting in the formation of a  
complex comprising the fusion protein bound to a receptor through the RP  
ligand and to the recombinant DNA vector through the DNA BP. An RP  
library (AAR91450-506) was screened with D32.39 and a six amino acid  
region of dynorphin B (RQFKV), an opioid peptide, was found to be the  
preferred recognition sequence for D32.39.  
(Updated on 25-MAR-2003 to correct PF field.)  
Sequence 12 AA;  
Query Match 62.2%; Score 28; DB 17; Length 12;  
Best Local Similarity 100.0%; Pred. No. 2e+02;  
Matches 4; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  
QY 2 VRWH 5  
DB |||||  
2 VRWH 5  
RESULT 43  
AAW25286  
ID AAW25286 standard; peptide; 12 AA.  
XX  
AC AAW25286;  
XX  
DT 14-OCT-1997 (first entry)  
XX  
DE Antibody D32.39 epitope #53.  
KW PCR; polymerase chain reaction; primer; amplify; lacI; headpiece domain;  
KW random peptide library; DNA binding protein; receptor ligand; dimer;  
KW fusion protein; epitope; antibody.  
XX  
OS Synthetic.

XX FH Key Location/Qualifiers  
FT Region 3..8  
FT /note= "D32.39 recognition site"  
XX  
XX PN WO9640987-A1.  
XX  
XX PD 19-DEC-1996.  
XX  
XX PF 07-JUN-1996; 96WO-US09809.  
XX  
XX PR 26-OCT-1995; 95US-0548540.  
XX PR 07-JUN-1995; 95US-0484090.  
XX  
XX PA (AFFY-) AFFYMAX TECHNOLOGIES NV.  
XX  
XX PI Cull MG, Gates CM, Miller JF, Schatz PJ, Stemmer WPC;  
XX  
XX DR WPI; 1997-087065/08.  
XX  
XX Random peptide library and affinity enrichment methods for screening  
PT it - useful to identify peptide(s) that bind receptor mols. of  
PT interest, useful for therapeutic, diagnostic and related purposes  
XX  
XX Example 4; Fig 3b; 149pp; English.  
XX  
XX AA25231-W25288 represent epitopes for the antibody D32.39. These  
CC sequences were isolated by a method of the invention to isolate a DNA  
CC binding protein, or a peptide with specific affinity for a receptor. The  
CC method comprises providing a recombinant DNA vector encoding a peptide  
CC having specific affinity for a receptor. A library of oligonucleotides  
CC encoding different potential DNA binding proteins is inserted in-frame  
CC into the vector to create a fusion protein library. Host cells are  
CC transformed, and cultured to express the fusion protein. If a fusion  
CC protein comprises a potential DNA binding protein with affinity for the  
CC vector, the fusion protein binds to the vector to form a complex. The  
CC host cells are lysed to isolate the complexes which are contacted with a  
CC receptor to induce peptide binding to the receptor. The random peptide  
CC library and the methods for screening it can be used to identify peptides  
CC that bind receptor molecules of interest. The peptides can be used for  
CC therapeutic, diagnostic and related purposes, e.g. to bind the receptor,  
CC or an analogue, and so inhibit or promote the activity of the receptor.  
CC The method of affinity enrichment allows a very large library of peptides  
CC to be screened, and by identifying the peptide de novo, the sequence or  
CC structure of the receptor molecule or the natural binding partner of the  
CC receptor need not be known.  
XX  
XX Sequence 12 AA;  
SQ  
Query Match 62.2%; Score 28; DB 18; Length 12;  
Best Local Similarity 100.0%; Pred. No. 2e+02;  
Matches 4; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  
QY 2 VRWH 5  
Db ||||  
2 VRWH 5  
RESULT 44  
AAB60032  
ID AAB60032 standard; Peptide; 12 AA.  
XX  
XX AC AAB60032;  
XX  
XX DT 05-NOV-2001 (first entry)  
XX  
XX DE Internalising peptide SEQ ID NO: 47.  
XX  
XX KW Internalising peptide; transport; apoptosis; arthritis; cancer;  
KW stem cell; cell differentiation; immune response stimulation;  
KW HIV vaccine.  
XX  
XX OS Synthetic.

XX WO200115511-A2.  
PN  
XX  
XX PD 08-MAR-2001.  
XX  
XX PF 31-AUG-2000; 2000WO-US24034.  
XX  
XX PR 01-SEP-1999; 99US-0151980.  
XX PR 13-MAR-2000; 2000US-0188944.  
XX  
XX PA (UYPI-) UNIV PITTSBURGH.  
XX  
XX PI Robbins PD, Mi Z, Frizzell R, Glorioso JC, Gambotto A;  
XX  
XX DR WPI; 2001-273309/28.  
XX  
XX PT Peptides that facilitate uptake and cytoplasmic and/or nuclear  
PT transport of proteins, DNA and viruses, useful, e.g. for facilitating  
PT uptake of antigens in immunogenic compositions -  
XX  
XX Claim 1; Page 123; 129pp; English.  
XX  
XX The present invention provides the sequences of 75 peptides which  
CC facilitate the uptake and transport of viruses, proteins and nucleic  
CC acids. These internalising peptides can be used for transport into the  
CC cytoplasm or the nucleus. They are useful for facilitating uptake into  
CC the cell, inducing apoptosis, for example in the treatment of arthritis  
CC and cancer, to expand a population of stem cells or differentiated cells,  
CC to stimulate cell differentiation, facilitate the integration of AAV into  
CC the genome of a cell, and to stimulate an immune response, for example in  
CC the case of a HIV vaccine. The present sequence is one of the peptides of  
CC the invention.  
XX  
XX SQ Sequence 12 AA;  
SQ  
Query Match 62.2%; Score 28; DB 22; Length 12;  
Best Local Similarity 60.0%; Pred. No. 2e+02;  
Matches 3; Conservative 0; Mismatches 2; Indels 0; Gaps 0;  
QY 1 WVRWH 5  
Db | ||  
3 WTPWH 7  
RESULT 45  
ABP46201  
ID ABP46201 standard; peptide; 13 AA.  
XX  
XX AC ABP46201;  
XX  
XX DT 19-AUG-2002 (first entry)  
XX  
XX DE Human BlyS binding scFv VH CDR3 SEQ ID 2212.  
XX  
XX KW BlyS; B lymphocyte stimulator; TNF superfamily; human; cytostatic;  
KW tumour necrosis factor; B cell proliferation; B cell differentiation;  
KW immunosuppressive; immunostimulant; immunomodulatory; antirheumatic;  
KW antiAIDS; vaccine; cancer; immune; autoimmune disorder; immunodeficiency;  
KW systemic lupus erythematosus; rheumatoid arthritis; CVID; AIDS;  
KW common variable immunodeficiency; acquired immunodeficiency syndrome.  
XX  
XX OS Homo sapiens.  
XX  
XX PN WO200202641-A1.  
XX  
XX PD 10-JAN-2002.  
XX  
XX PF 15-JUN-2001; 2001WO-US19110.  
XX  
XX PR 16-JUN-2000; 2000US-212210P.  
XX PR 17-OCT-2000; 2000US-240816P.  
XX PR 16-MAR-2001; 2001US-276248P.  
XX PR 21-MAR-2001; 2001US-277379P.

PR 25-MAY-2001; 2001US-293499P.  
XX  
PA (HUMA-) HUMAN GENOME SCI INC.  
PA (CAMB-) CAMBRIDGE ANTIBODY TECHNOLOGY.  
XX  
PI Ruben SM, Barash SC, Choi GH, Vaughan T, Hilbert D;  
XX  
DR WPI; 2002-114799/15.  
XX  
XX  
PT Antibodies against B Lymphocyte Stimulating polypeptides, useful for  
PT the diagnosis and treatment of cancers and immune disorders -  
XX  
PS Claim 2; Page 2952; 3148pp; English.  
XX  
CC This invention describes novel antibodies that immunospecifically bind to  
CC B Lymphocyte Stimulator (BLYS) polypeptides. BLYS is a member of the  
CC tumour necrosis factor (TNF) super family and induces B cell  
CC proliferation and differentiation. The antibodies of the invention have  
CC cytostatic, immunosuppressive, immunostimulant, immunomodulatory,  
CC antirheumatic and antiAIDS activity and can be used in vaccines to  
CC inhibit the expression and activity of BLYS. The antibodies bind to BLYS  
CC and so may be used to detect and quantitate the presence of BLYS in  
CC biological samples and may be used in this way to diagnose disease  
CC associated with aberrant expression of BLYS. They may also be  
CC administered to treat diseases associated with aberrant BLYS expression  
CC and activity such as cancer, immune, and autoimmune disorders and  
CC diseases, e.g. systemic lupus erythematosus, rheumatoid arthritis,  
CC immunodeficiency (e.g. common variable immunodeficiency (CVID) and  
CC acquired immunodeficiency syndrome (AIDS)). ABP43990-ABP47228 represent  
CC the antibodies and fragments of the antibodies described in the method  
CC of the invention.  
XX  
SQ Sequence 13 AA;  
  
Query Match 62.2%; Score 28; DB 23; Length 13;  
Best Local Similarity 50.0%; Pred. No. 2.2e+02;  
Matches 3; Conservative 1; Mismatches 2; Indels 0; Gaps 0;  
  
QY 1 WVRWHF 6  
| | | | |  
Db 6 WPNWYF 11

Search completed: December 12, 2003, 10:29:02  
Job time : 31.3 secs

GenCore version 5.1.6  
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OM protein - protein search, using sw model

Run on: December 3, 2003, 11:44:55 ; Search time 7.33333 Seconds  
(without alignments)  
38.476 Million cell updates/sec

Title: US-09-912-414-2  
Perfect score: 45  
Sequence: 1 WVRWHF 6

Scoring table: BLOSUM62  
Gapop 10.0 , Gapext 0.5

Searched: 127863 seqs, 47026705 residues

Total number of hits satisfying chosen parameters: 795

Minimum DB seq length: 0  
Maximum DB seq length: 15

Post-processing: Minimum Match 0%  
Maximum Match 100%  
Listing first 45 summaries

Database : SwissProt\_41:\*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	DB ID	Description
1	22	48.9	9	1 LITR_PHYRO	P08946 phyllomedusa
2	22	48.9	11	1 RANC_RANPI	P08951 rana pipien
3	20	44.4	9	1 LITO_LITAU	P08945 litoria aur
4	20	44.4	13	1 BOML_PSEGU	P42991 pseudophryn
5	19.5	43.3	5	1 UF01_MOUSE	P38639 mus musculus
6	19	42.2	10	1 LABA_JATMU	P13270 jatropha mu
7	18	40.0	9	1 COW_CONVE	P83047 conus ventr
8	18	40.0	13	1 YPNP_PHOLU	P41122 photorhabdu
9	17	37.8	11	1 MLG_THETS	P41989 theromyzon
10	17	37.8	13	1 EI21_LITRU	P82097 litoria rub
11	17	37.8	13	1 EI22_LITRU	P82098 litoria rub
12	17	37.8	13	1 TEML_RANTE	P57104 rana tempor
13	16	35.6	8	1 ACI_THUAL	P18691 thunnus alb
14	16	35.6	13	1 MLA_ANOCA	P41589 anolis caro
15	16	35.6	13	1 MLA_CAMDR	P01198 camelus dro
16	16	35.6	14	1 LPW_RHIME	P18854 rhizobium m
17	15	33.3	7	1 TPFY_PACDA	P83455 pachymedusa
18	15	33.3	10	1 AEGL_AGRAE	P83465 agrocybe ae
19	15	33.3	12	1 RF1_CONSP	P58805 conus spuri
20	15	33.3	15	1 CX3B_CONQU	P58842 conus querc
21	15	33.3	15	1 GLN2_PINPS	P81107 pinus pinas
22	14	31.1	10	1 BPP2_BOTJA	P01022 bothrops ja
23	14	31.1	10	1 FARP_MYTED	P42560 mytilus edu
24	14	31.1	10	1 GRP_RANRI	P23260 rana ridibu
25	14	31.1	11	1 CA22_LITCI	P82088 litoria cit
26	14	31.1	11	1 CA42_LITCI	P82092 litoria cit
27	14	31.1	13	1 BPP1_BOTJA	P01020 bothrops ja
28	14	31.1	13	1 CXA2_CONGE	P01520 conus geogr
29	14	31.1	14	1 ALYT_ALYOB	P08944 alytes obst
30	14	31.1	14	1 MAST_PARID	P42716 parapolylbia
31	14	31.1	14	1 MAST_VESBA	P21654 vespa basal
32	14	31.1	14	1 MAST_VESXA	P01515 vespa xanth
33	14	31.1	15	1 AH2_PRUSE	P29260 prunus sero

34	14	31.1	15	1 DCMN_PSECH	P19917 pseudomonas
35	14	31.1	15	1 MK2A_PALPR	P80409 palomena pr
36	14	31.1	15	1 RM12_YEAST	P36522 saccharomyc
37	13	28.9	5	1 BPP7_BOTIN	P30425 bothrops in
38	13	28.9	9	1 NEF_HV1Z8	P12481 human immun
39	13	28.9	10	1 APE_CAPGI	P80474 capnocytoph
40	13	28.9	10	1 GON1_ALLMI	P37041 alligator m
41	13	28.9	10	1 GON2_CHEPR	P80678 chelyosoma
42	13	28.9	10	1 GON2_CHICK	P37043 gallus gall
43	13	28.9	10	1 GON3_ONCKE	P20367 oncorhynchu
44	13	28.9	10	1 NO40_TOBAC	P55962 nicotiana t
45	13	28.9	12	1 UP01_CAEEL	P55954 caenorhabdi

ALIGNMENTS

RESULT 1  
LITR\_PHYRO  
ID LITR\_PHYRO STANDARD; PRT; 9 AA.  
AC P08946;  
DT 01-NOV-1988 (Rel. 09, Created)  
DT 01-FEB-1994 (Rel. 28, Last sequence update)  
DT 15-SEP-2003 (Rel. 42, Last annotation update)  
DE Rhodei-litorin.  
OS Phyllomedusa rohdei (Rohde's leaf frog).  
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
OC Amphibia; Batrachia; Anura; Neobatrachia; Bufonoidea; Hylidae;  
OC Phyllomedusinae; Phyllomedusa.  
OX NCBI\_TaxID=8394;  
RN [1]

RP SEQUENCE.  
RC TISSUE=Skin secretion;  
RX MEDLINE=85127560; PubMed=3838283;  
RA Barra D., Erspamer G.F., Simmaco M., Bossa F., Melchiorri P.,  
RA Erspamer V.;  
RT "Rohdei-litorin: a new peptide from the skin of Phyllomedusa rohdei.";  
RL FEBS Lett. 182:53-56(1985).  
CC -!- SUBCELLULAR LOCATION: Secreted.  
CC -!- TISSUE SPECIFICITY: Skin.  
CC -!- SIMILARITY: BELONGS TO THE BOMBESIN/NEUROMEDIN B/RANATENSIN  
CC FAMILY.  
CC PIR; S07241; S07241.  
DR InterPro: IPR000874; Bombesin.  
DR Pfam; PF02044; Bombesin; 1.  
DR PROSITE; PS00257; BOMBESIN; 1.  
KW Amphibian defense peptide; Bombesin family; Amidation;  
KW Pyrrolidone carboxylic acid.  
FT MOD\_RES 1 1 PYRROLIDONE CARBOXYLIC ACID.  
FT MOD\_RES 9 9 AMIDATION.  
SQ SEQUENCE 9 AA; 1090 MW; 4ECCC1E861ADC377 CRC64;

Query Match 48.9%; Score 22; DB 1; Length 9;  
Best Local Similarity 50.0%; Pred. No. 1.3e+05;  
Matches 3; Conservative 0; Mismatches 3; Indels 0; Gaps 0;

QY 1 WVRWHF 6  
Db 3 WATGHF 8

RESULT 2  
RANC\_RANPI  
ID RANC\_RANPI STANDARD; PRT; 11 AA.  
AC P08951;  
DT 01-NOV-1988 (Rel. 09, Created)  
DT 01-NOV-1988 (Rel. 09, Last sequence update)  
DT 15-SEP-2003 (Rel. 42, Last annotation update)  
DE Ranatensin-C.  
OS Rana pipiens (Northern leopard frog).  
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
OC Amphibia; Batrachia; Anura; Neobatrachia; Ranioidea; Rana.  
OX NCBI\_TaxID=8404;



[1]  
SEQUENCE.  
RC TISSUE=Skin secretion;  
RX MEDLINE=84131098; PubMed=6141890;  
RA Nakajima T.;  
RL Unpublished results, cited by:  
RL Erspamer V., Erspamer G.F., Mazzanti G., Endean R.;  
RL Comp. Biochem. Physiol. 77C:99-108(1984).  
CC -!- SUBCELLULAR LOCATION: Secreted.  
CC -!- TISSUE SPECIFICITY: Skin.  
CC -!- SIMILARITY: BELONGS TO THE BOMBESIN/NEUROMEDIN B/RANATENSIN  
CC FAMILY.  
DR InterPro: IPR000874; Bombesin.  
DR Pfam: PF02044; Bombesin; 1.  
DR PROSITE: PS00257; BOMBESIN; 1.  
KW Amphibian defense peptide; Bombesin family; Amidation.  
FT MOD\_RES 11 AMIDATION.  
SQ SEQUENCE 11 AA; 1304 MW; D6C9885A61ADC366 CRC64;  
  
Query Match 48.9%; Score 22; DB 1; Length 11;  
Best Local Similarity 50.0%; Pred. No. 2e+02;  
Matches 3; Conservative 0; Mismatches 3; Indels 0; Gaps 0;  
  
QY 1 WVRWHF 6  
| |  
Db 5 WATGHF 10  
  
RESULT 3  
LITO\_LITAU STANDARD; PRT; 9 AA.  
ID LITO\_LITAU  
AC P08945;  
DT 01-NOV-1988 (Rel. 09, Created)  
DT 01-FEB-1994 (Rel. 28, Last sequence update)  
DT 15-SEP-2003 (Rel. 42, Last annotation update)  
DE Litorin.  
OS Litoria aurea (Green and golden bell frog).  
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
OC Amphibia; Batrachia; Anura; Neobatrachia; Bufonoidea; Hylidae;  
OC Pelodyadinae; Litoria.  
OX NCBI\_TaxID=8371;  
RN [1]  
RP SEQUENCE.  
RC TISSUE=Skin secretion;  
RX MEDLINE=75187011; PubMed=1140241;  
RA Anastasi A., Erspamer V., Endean R.;  
RT "Aminoacid composition and sequence of litorin, a bombesin-like nonapeptide from the skin of the Australian leptodactylid frog Litoria aurea.";  
RL Experientia 31:510-511(1975).  
RN [2]  
RP SEQUENCE (METHYLATED VARIANT).  
RC TISSUE=Skin secretion;  
RX MEDLINE=78003546; PubMed=908397;  
RA Anastasi A., Montecucchi P.C., Angelucci F., Erspamer V., Endean R.;  
RT "Glu(OMe)3-litorin, the second bombesin-like peptide occurring in methanol extracts of the skin of the Australian frog Litoria aurea.";  
RL Experientia 33:1289-1289(1977).  
CC -!- SUBCELLULAR LOCATION: Secreted.  
CC -!- TISSUE SPECIFICITY: Skin.  
CC -!- SIMILARITY: BELONGS TO THE BOMBESIN/NEUROMEDIN B/RANATENSIN  
CC FAMILY.  
DR PIR: S07204; S07204.  
DR InterPro: IPR000874; Bombesin.  
DR Pfam: PF02044; Bombesin; 1.  
DR PROSITE: PS00257; BOMBESIN; 1.  
KW Amphibian defense peptide; Bombesin family; Amidation; Methylation;  
KW Pyrrolidone carboxylic acid.  
FT MOD\_RES 1 1 PYRROLIDONE CARBOXYLIC ACID.  
FT MOD\_RES 2 2 METHYLATION (PARTIAL).  
FT MOD\_RES 9 9 AMIDATION.  
SQ SEQUENCE 9 AA; 1103 MW; D7CCCL862CDC366 CRC64;

Query Match 44.4%; Score 20; DB 1; Length 9;  
Best Local Similarity 50.0%; Pred. No. 1.3e+05;  
Matches 3; Conservative 0; Mismatches 3; Indels 0; Gaps 0;  
  
QY 1 WVRWHF 6  
| |  
Db 3 WATGHF 8  
  
RESULT 4  
BOML\_PSEGU STANDARD; PRT; 13 AA.  
ID BOML\_PSEGU  
AC P42991;  
DT 01-NOV-1995 (Rel. 32, Created)  
DT 01-NOV-1995 (Rel. 32, Last sequence update)  
DT 15-SEP-2003 (Rel. 42, Last annotation update)  
DE Bombesin-like peptide L (PG-L).  
OS Pseudophryne guentheri (Guenther's toadlet).  
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
OC Amphibia; Batrachia; Anura; Neobatrachia; Bufonoidea; Myobatrachidae;  
OC Myobatrachinae; Pseudophryne.  
OX NCBI\_TaxID=30349;  
RN [1]  
RP SEQUENCE.  
RC TISSUE=Skin secretion;  
RX MEDLINE=90287814; PubMed=2356157;  
RA Simmaco M., Severini C., de Biase D., Barra D., Bossa F.,  
RA Roberts J.D., Melchiorri P., Erspamer V.;  
RT "Six novel tachykinin- and bombesin-related peptides from the skin of the Australian frog Pseudophryne guentheri.";  
RL Peptides 11:299-304(1990).  
CC -!- SUBCELLULAR LOCATION: Secreted.  
CC -!- TISSUE SPECIFICITY: Skin.  
CC -!- SIMILARITY: BELONGS TO THE BOMBESIN/NEUROMEDIN B/RANATENSIN  
CC FAMILY.  
DR PIR: A60409; A60409.  
DR InterPro: IPR000874; Bombesin.  
DR Pfam: PF02044; Bombesin; 1.  
DR PROSITE: PS00257; BOMBESIN; 1.  
KW Amphibian defense peptide; Bombesin family; Amidation;  
KW Pyrrolidone carboxylic acid.  
FT MOD\_RES 1 1 PYRROLIDONE CARBOXYLIC ACID.  
FT MOD\_RES 13 13 AMIDATION.  
SQ SEQUENCE 13 AA; 1372 MW; D6DE0D24BD98C366 CRC64;  
  
Query Match 44.4%; Score 20; DB 1; Length 13;  
Best Local Similarity 50.0%; Pred. No. 4.9e+02;  
Matches 3; Conservative 0; Mismatches 3; Indels 0; Gaps 0;  
  
QY 1 WVRWHF 6  
| |  
Db 7 WATGHF 12  
  
RESULT 5  
UF01\_MOUSE STANDARD; PRT; 5 AA.  
ID UF01\_MOUSE  
AC P38639;  
DT 01-OCT-1994 (Rel. 30, Created)  
DT 01-OCT-1994 (Rel. 30, Last sequence update)  
DT 01-FEB-1995 (Rel. 31, Last annotation update)  
DE Unknown protein from 2D-page of fibroblasts (P19) (Fragment).  
OS Mus musculus (Mouse).  
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
OC Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.  
OX NCBI\_TaxID=10090;  
RN [1]  
RP SEQUENCE.  
RC TISSUE=Fibroblast;  
RX MEDLINE=95009907; PubMed=7523108;  
RA Merrick B.A., Patterson R.M., Wichter L.L., He C., Selkirk J.K.;  
RT "Separation and sequencing of familial and novel murine proteins using preparative two-dimensional gel electrophoresis.";

RL Electrophoresis 15:735-745(1994).  
CC -!- MISCELLANEOUS: ON THE 2D-GEL THE DETERMINED PI OF THIS UNKNOWN  
CC PROTEIN IS: 6.6, ITS MW IS: 19 kDa.  
FT NON\_TER 5 5  
SQ SEQUENCE 5 AA; 717 MW; 7364087043100000 CRC64;

Query Match 43.3%; Score 19.5; DB 1; Length 5;  
Best Local Similarity 60.0%; Pred. No. 1.3e+05;  
Matches 3; Conservative 1; Mismatches 0; Indels 1; Gaps 1;

QY 1 WVRW 4  
|:|  
Db 1 WIGRW 5

RESULT 6  
LABA\_JATMU  
ID LABA\_JATMU STANDARD; PRT; 10 AA.  
AC P13270;  
DT 01-JAN-1990 (Rel. 13, Created)  
DT 01-JAN-1990 (Rel. 13, Last sequence update)  
DT 28-FEB-2003 (Rel. 41, Last annotation update)  
DE Labaditin.  
OS Jatropa multifida (Physic nut).  
OC Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;  
OC Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; Rosidae;  
OC eurosids I; Malpighiales; Euphorbiaceae; Jatropa.  
OX NCBI\_TaxID=3996;  
RN [1]  
RP SEQUENCE.  
RC TISSUE=Latex;  
RA Kosasi S., van der Sluis W.G., Boelens R., T'Hart L.A., Labadie R.P.;  
RT "Labaditin, a novel cyclic decapeptide from the latex of Jatropa  
RT multifida L. (Euphorbiaceae). Isolation and sequence determination  
RT by means of two-dimensional NMR.";  
RL FEBS Lett. 256:91-96(1989).  
CC -!- FUNCTION: LABADITIN IS AN ACTIVE PEPTIDE WHICH INHIBITS THE  
CC CLASSICAL PATHWAY OF COMPLEMENT ACTIVATION IN VITRO. ACTIVITY  
CC SEEMS TO BE BASED ON AN INTERACTION WITH C1.  
CC -!- PTM: This is a cyclic peptide.  
CC -!- DISEASE: LATEX OF THIS PLANT IS USED IN FOLKLORIC MEDICINE FOR  
CC TREATMENT OF INFECTED WOUNDS, SKINS INFECTIONS AND SCABIES.  
SQ SEQUENCE 10 AA; 1089 MW; D98AAD6362D1B362 CRC64;

Query Match 42.2%; Score 19; DB 1; Length 10;  
Best Local Similarity 50.0%; Pred. No. 5.6e+02;  
Matches 2; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

QY 1 WVRW 4  
|:|  
Db 4 WTVW 7

RESULT 7  
COW\_CONVE  
ID COW\_CONVE STANDARD; PRT; 9 AA.  
AC P83047;  
DT 16-OCT-2001 (Rel. 40, Created)  
DT 16-OCT-2001 (Rel. 40, Last sequence update)  
DT 28-FEB-2003 (Rel. 41, Last annotation update)  
DE Contryphan-Vn.  
OS Conus ventricosus (Mediterranean cone).  
OC Eukaryota; Metazoa; Mollusca; Gastropoda; Orthogastropoda;  
OC Apogastropoda; Caenogastropoda; Sorbeoconcha; Hypsogastropoda;  
OC Neogastropoda; Conoidea; Conidae; Conus.  
OX NCBI\_TaxID=117992;  
RN [1]  
RP SEQUENCE, SYNTHESIS, AND MASS SPECTROMETRY.  
RC TISSUE=Venom;  
RX MEDLINE=21547785; PubMed=11688995;  
RA Massilia G.R., Schinina M.E., Ascenzi P., Politicelli F.;  
RT "Contryphan-Vn: a novel peptide from the venom of the Mediterranean  
RT snail Conus ventricosus.";

RL Biochem. Biophys. Res. Commun. 288:908-913(2001).  
CC -!- SUBCELLULAR LOCATION: Secreted.  
CC -!- TISSUE SPECIFICITY: Expressed by the venom duct.  
CC -!- MASS SPECTROMETRY: MW=1088.6; METHOD=MALDI.  
CC -!- SIMILARITY: BELONGS TO THE CONTRYPHAN FAMILY.  
KW Toxin; Amidation; D-amino acid.  
FT DISULFID 3 9  
FT MOD\_RES 5 5 D-TRYPTOPHAN.  
FT MOD\_RES 9 9 AMIDATION.  
SQ SEQUENCE 9 AA; 1091 MW; 8D38676323676EBA CRC64;

Query Match 40.0%; Score 18; DB 1; Length 9;  
Best Local Similarity 50.0%; Pred. No. 1.3e+05;  
Matches 2; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

QY 1 WVRW 4  
|:|  
Db 5 WKPW 8

RESULT 8  
YPNP\_PHOLU  
ID YPNP\_PHOLU STANDARD; PRT; 13 AA.  
AC P41122;  
DT 01-FEB-1995 (Rel. 31, Created)  
DT 01-FEB-1995 (Rel. 31, Last sequence update)  
DT 16-OCT-2001 (Rel. 40, Last annotation update)  
DE Hypothetical protein in pnp 3'region (ORF3) (Fragment).  
OS Photorhabdus luminescens (Xenorhabdus luminescens).  
OC Bacteria; Proteobacteria; Gammaproteobacteria; Enterobacteriales;  
OC Enterobacteriaceae; Photorhabdus.  
OX NCBI\_TaxID=29488;  
RN [1]  
RP SEQUENCE FROM N.A.  
RC STRAIN=K122;  
RX MEDLINE=94266731; PubMed=8206856;  
RA Clarke D.J., Dowds B.C.A.;  
RT "The gene coding for polynucleotide phosphorylase in Photorhabdus sp.  
RT strain K122 is induced at low temperatures.";  
RL J. Bacteriol. 176:3775-3784(1994).  
CC -----  
CC This SWISS-PROT entry is copyright. It is produced through a collaboration  
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CC or send an email to [license@isb-sib.ch](mailto:license@isb-sib.ch)).  
CC -----  
CC EMBL; X76069; CAAS3672.1; -.  
KW Hypothetical protein.  
FT NON\_TER 13 13  
SQ SEQUENCE 13 AA; 1634 MW; 64774A4F6267A364 CRC64;

Query Match 40.0%; Score 18; DB 1; Length 13;  
Best Local Similarity 50.0%; Pred. No. 1.1e+03;  
Matches 2; Conservative 2; Mismatches 0; Indels 0; Gaps 0;

QY 1 WVRW 4  
|:|  
Db 4 FLRW 7

RESULT 9  
MLG\_THETS  
ID MLG\_THETS STANDARD; PRT; 11 AA.  
AC P41989;  
DT 01-NOV-1995 (Rel. 32, Created)  
DT 01-NOV-1995 (Rel. 32, Last sequence update)  
DT 16-OCT-2001 (Rel. 40, Last annotation update)  
DE Melanotropin gamma (Gamma-melanocyte stimulating hormone) (Gamma-MSH).  
OS Theromyzon tessulatum (Leech).  
OC Eukaryota; Metazoa; Annelida; Clitellata; Hirudinida; Hirudinea;

OC Rhynchobdellida; Glossiphoniidae; Theromyzon.  
OX NCBI\_TaxID=13286;  
RN [1]  
RP SEQUENCE.  
RC TISSUE=Brain;  
RX MEDLINE=94298944; PubMed=8026574;  
RA Salzet M., Watzet C., Bulet P., Malecha J.;  
RT "Isolation and structural characterization of a novel peptide related  
to gamma-melanocyte stimulating hormone from the brain of the leech  
RT Theromyzon tessellatum";  
RL FEBS Lett. 348:102-106(1994).  
CC -!- SIMILARITY: BELONGS TO THE POMC FAMILY.  
DR PIR; S45698; S45698.  
KW Hormone; Amidation.  
FT MOD RES 11 11 AMIDATION.  
SQ SEQUENCE 11 AA; 1486 MW; 2DB8FACE6409C1E8 CRC64;  
  
Query Match 37.8%; Score 17; DB 1; Length 11;  
Best Local Similarity 50.0%; Pred. No. 1.3e+03;  
Matches 3; Conservative 1; Mismatches 2; Indels 0; Gaps 0;  
  
QY 1 WVRWHF 6  
Db 1 YVMGHF 6

RESULT 10  
EI21\_LITRU  
ID EI21\_LITRU STANDARD; PRT; 13 AA.  
AC P82097;  
DT 28-FEB-2003 (Rel. 41, Created)  
DT 28-FEB-2003 (Rel. 41, Last sequence update)  
DT 15-SEP-2003 (Rel. 42, Last annotation update)  
DE Electrin 2.1.  
OS Litoria rubella (Desert tree frog).  
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
OC Amphibia; Batrachia; Anura; Neobatrachia; Bufonoidea; Hylidae;  
OC Pelodyadinae; Litoria.  
OX NCBI\_TaxID=104895;  
RN [1]  
RP SEQUENCE.  
RC TISSUE=Skin secretion;  
RA Wabnitz P.A., Bowie J.H., Tyler M.J., Wallace J.C.;  
RT "Peptides from the skin glands of the Australian buzzing tree frog  
RT Litoria electrica. Comparison with the skin peptides from Litoria  
RT rubella";  
RL Aust. J. Chem. 52:639-645(1999).  
CC -!- SUBCELLULAR LOCATION: Secreted.  
CC -!- TISSUE SPECIFICITY: Skin.  
KW Amphibian defense peptide; Amidation.  
FT MOD RES 13 13 AMIDATION.  
SQ SEQUENCE 13 AA; 1599 MW; C1808EF326F57322 CRC64;  
  
Query Match 37.8%; Score 17; DB 1; Length 13;  
Best Local Similarity 66.7%; Pred. No. 1.5e+03;  
Matches 2; Conservative 1; Mismatches 0; Indels 0; Gaps 0;  
  
QY 2 VRW 4  
Db 6 VKW 8

RESULT 11  
EI22\_LITRU  
ID EI22\_LITRU STANDARD; PRT; 13 AA.  
AC P82098;  
DT 28-FEB-2003 (Rel. 41, Created)  
DT 28-FEB-2003 (Rel. 41, Last sequence update)  
DT 15-SEP-2003 (Rel. 42, Last annotation update)  
DE Electrin 2.2.  
OS Litoria rubella (Desert tree frog).  
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
OC Amphibia; Batrachia; Anura; Neobatrachia; Bufonoidea; Hylidae;

OC Pelodyadinae; Litoria.  
OX NCBI\_TaxID=104895;  
RN [1]  
RP SEQUENCE.  
RC TISSUE=Skin secretion;  
RA Wabnitz P.A., Bowie J.H., Tyler M.J., Wallace J.C.;  
RT "Peptides from the skin glands of the Australian buzzing tree frog  
RT Litoria electrica. Comparison with the skin peptides from Litoria  
RT rubella";  
RL Aust. J. Chem. 52:639-645(1999).  
CC -!- SUBCELLULAR LOCATION: Secreted.  
CC -!- TISSUE SPECIFICITY: Skin.  
KW Amphibian defense peptide; Amidation.  
FT MOD RES 13 13 AMIDATION.  
SQ SEQUENCE 13 AA; 1598 MW; C1808EF33B357322 CRC64;  
  
Query Match 37.8%; Score 17; DB 1; Length 13;  
Best Local Similarity 66.7%; Pred. No. 1.5e+03;  
Matches 2; Conservative 1; Mismatches 0; Indels 0; Gaps 0;  
  
QY 2 VRW 4  
Db 6 VKW 8

RESULT 12  
TEML\_RANTE  
ID TEML\_RANTE STANDARD; PRT; 13 AA.  
AC P57104;  
DT 16-OCT-2001 (Rel. 40, Created)  
DT 16-OCT-2001 (Rel. 40, Last sequence update)  
DT 15-SEP-2003 (Rel. 42, Last annotation update)  
DE Temporin L.  
OS Rana temporaria (European common frog).  
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
OC Amphibia; Batrachia; Anura; Neobatrachia; Ranoidea; Ranidae; Rana.  
OX NCBI\_TaxID=8407;  
RN [1]  
RP SEQUENCE.  
RC TISSUE=Skin secretion;  
RX MEDLINE=97175050; PubMed=9022710;  
RA Simmaco M., Mignogna G., Canofeni S., Miele R., Mangoni M.L.,  
RA Barra D.;  
RT "Temporins, antimicrobial peptides from the European red frog Rana  
RT temporaria";  
RL Eur. J. Biochem. 242:788-792(1996).  
CC -!- FUNCTION: HAS ANTIBACTERIAL ACTIVITY AGAINST GRAM-NEGATIVE AND  
CC GRAM-POSITIVE BACTERIA.  
CC -!- SUBCELLULAR LOCATION: Secreted.  
CC -!- TISSUE SPECIFICITY: Skin.  
CC -!- SIMILARITY: Belongs to the brevinin family.  
KW Amphibian defense peptide; Antibiotic; Amidation.  
FT MOD RES 13 13 AMIDATION.  
SQ SEQUENCE 13 AA; 1641 MW; 9EBDCB1FAFF7C325 CRC64;  
  
Query Match 37.8%; Score 17; DB 1; Length 13;  
Best Local Similarity 50.0%; Pred. No. 1.5e+03;  
Matches 2; Conservative 2; Mismatches 0; Indels 0; Gaps 0;  
  
QY 1 WVRW 4  
Db 1 FVOW 4

RESULT 13  
ACI\_THUAL  
ID ACI\_THUAL STANDARD; PRT; 8 AA.  
AC P18691;  
DT 01-NOV-1990 (Rel. 16, Created)  
DT 01-NOV-1990 (Rel. 16, Last sequence update)  
DT 01-NOV-1990 (Rel. 16, Last annotation update)  
DE Angiotensin-converting enzyme inhibitor.  
OS Thunnus albacares (Yellowfin tuna) (Neothunnus macropterus).

OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
OC Actinopterygii; Neopterygii; Teleostei; Euteleostei; Neoteleostei;  
OC Acanthomorpha; Acanthopterygii; Percomorpha; Perciformes; Scombroidei;  
OC Scombridae; Thunnus.  
OX NCBI\_TaxID=8236;  
RN [1]  
RP SEQUENCE.  
RC TISSUE=Muscle;  
RX MEDLINE=88326322; PubMed=3415688;  
RA Kohama Y., Matsumoto S., Oka H., Teramoto T., Okabe M., Mimura T.;  
RT "Isolation of angiotensin-converting enzyme inhibitor from tuna  
muscle.";  
RL Biochem. Biophys. Res. Commun. 155:332-337(1988).  
DR PIR; A31570; A31570.  
SQ SEQUENCE 8 AA; 953 MW; 6AA863733051F1B7 CRC64;  
  
Query Match 35.6%; Score 16; DB 1; Length 8;  
Best Local Similarity 33.3%; Pred. No. 1.3e+05;  
Matches 1; Conservative 2; Mismatches 0; Indels 0; Gaps 0;  
  
Qy 2 VRW 4  
Db 4 IKW 6  
  
RESULT 14  
MLA\_ANOCA STANDARD; PRT; 13 AA.  
ID MLA\_ANOCA  
AC P41589;  
DT 01-NOV-1995 (Rel. 32, Created)  
DT 01-NOV-1995 (Rel. 32, Last sequence update)  
DT 16-OCT-2001 (Rel. 40, Last annotation update)  
DE Melanotropin alpha (Alpha-MSH).  
OS Anolis carolinensis (Green anole) (American chameleon).  
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
OC Lepidosauria; Squamata; Iguania; Iguanidae; Polychrotinae; Anolis.  
OX NCBI\_TaxID=28377;  
RN [1]  
RP SEQUENCE.  
RC TISSUE=Pituitary;  
RX MEDLINE=92270473; PubMed=1667689;  
RA Does R.M., Lancha A., Rand-Weaver M., Jankelow L., Adamczyk D.L.;  
RT "Detection of a novel sequence change in the major form of alpha-MSH  
isolated from the intermediate pituitary of the reptile, Anolis  
carolinensis.";  
RL Peptides 12:1261-1266(1991).  
CC -!- SIMILARITY: BELONGS TO THE POMC FAMILY.  
DR InterPro; IPR001941; Mcortin ACTH.  
DR Pfam; PF00976; ACTH\_domain; 1.  
KW Hormone; Amidation.  
FT MOD\_RES 13 13 AMIDATION.  
SQ SEQUENCE 13 AA; 1608 MW; FF990A7358BB09C1 CRC64;  
  
Query Match 35.6%; Score 16; DB 1; Length 13;  
Best Local Similarity 100.0%; Pred. No. 2.3e+03;  
Matches 2; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  
  
Qy 3 RW 4  
Db 8 RW 9  
  
RESULT 15  
MLA\_CAMDR STANDARD; PRT; 13 AA.  
ID MLA\_CAMDR  
AC P01198;  
DT 21-JUL-1986 (Rel. 01, Created)  
DT 21-JUL-1986 (Rel. 01, Last sequence update)  
DT 16-OCT-2001 (Rel. 40, Last annotation update)  
DE Melanotropin alpha (Alpha-MSH).  
OS Camelus dromedarius (Dromedary) (Arabian camel), and  
Equus caballus (Horse).  
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;

OC Mammalia; Eutheria; Cetartiodactyla; Tylopoda; Camelidae; Camelus.  
OX NCBI\_TaxID=9838, 9796;  
RN [1]  
RP SEQUENCE.  
RC SPECIES=C.dromedarius;  
RX MEDLINE=75146434; PubMed=1125179;  
RA Li C.H., Danho W.O., Chung D., Rao A.J.;  
RT "Isolation, characterization, and amino acid sequence of  
melanotropins from camel pituitary glands.";  
RL Biochemistry 14:947-952(1975).  
RN [2]  
RP SEQUENCE.  
RC SPECIES=Horse; TISSUE=Pituitary;  
RA Dixon J.S., Li C.H.;  
RT "The isolation and structure of alpha-melanocyte-stimulating hormone  
from horse pituitaries.";  
RL J. Am. Chem. Soc. 82:4568-4572(1960).  
CC -!- SIMILARITY: BELONGS TO THE POMC FAMILY.  
DR PIR; A01464; MTCMAD.  
DR PIR; A91785; MTHOAD.  
DR InterPro; IPR001941; Mcortin ACTH.  
DR Pfam; PF00976; ACTH\_domain; 1.  
KW Hormone; Acetylation; Amidation.  
FT MOD\_RES 1 1 ACETYLATION (IN ABOUT 50% OF CAMEL  
MOLECULES).  
FT MOD\_RES 13 13 AMIDATION.  
SQ SEQUENCE 13 AA; 1624 MW; FF991CA958BB09C1 CRC64;  
  
Query Match 35.6%; Score 16; DB 1; Length 13;  
Best Local Similarity 100.0%; Pred. No. 2.3e+03;  
Matches 2; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  
  
Qy 3 RW 4  
Db 8 RW 9  
  
Search completed: December 3, 2003, 11:51:51  
Job time : 9.33333 secs

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OM protein - protein search, using sw model

Run on: December 3, 2003, 11:48:05 ; Search time 26.3333 Seconds  
(without alignments)  
58.797 Million cell updates/sec

Title: US-09-912-414-9  
Perfect score: 31  
Sequence: 1 WXXWXP 6

Scoring table: BLOSUM62  
Gapop 10.0 , Gapext 0.5

Searched: 830525 seqs, 258052604 residues

Total number of hits satisfying chosen parameters: 3526

Minimum DB seq length: 0  
Maximum DB seq length: 15

Post-processing: Minimum Match 0%  
Maximum Match 100%  
Listing first 45 summaries

Database : SPTREMBL\_23:\*

- 1: sp\_archaea:\*
- 2: sp\_bacteria:\*
- 3: sp\_fungi:\*
- 4: sp\_human:\*
- 5: sp\_invertebrate:\*
- 6: sp\_mammal:\*
- 7: sp\_mhc:\*
- 8: sp\_organelle:\*
- 9: sp\_phage:\*
- 10: sp\_plant:\*
- 11: sp\_rodent:\*
- 12: sp\_virus:\*
- 13: sp\_vertibrate:\*
- 14: sp\_unclassified:\*
- 15: sp\_rvirus:\*
- 16: sp\_bacteriap:\*
- 17: sp\_archaeap:\*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	DB ID	Description
1	21	67.7	9	2 Q9R5M1	Q9r5m1 staphylococ
2	21	67.7	9	9 Q38366	Q38366 bacterioph
3	20	64.5	9	8 Q8SHF0	Q8shf0 chamealeo n
4	20	64.5	12	7 Q77919	Q77919 pseudotroph
5	20	64.5	13	4 Q16406	Q16406 homo sapien
6	20	64.5	15	2 Q53580	Q53580 rhodobacter
7	17	54.8	8	8 Q94VC1	Q94vc1 varanus rud
8	17	54.8	11	8 Q94V77	Q94v77 heloderma s
9	17	54.8	13	4 Q9UDC6	Q9udc6 homo sapien
10	17	54.8	14	10 Q9SAP8	Q9sap8 pisum sativ
11	16	51.6	8	8 Q94VF6	Q94vf6 varanus job
12	16	51.6	8	8 Q8WGD7	Q8wgd7 lomis hirta
13	16	51.6	8	8 Q94V88	Q94v88 varanus tri
14	16	51.6	8	8 Q9TD02	Q9td02 terranatos
15	16	51.6	8	8 Q9T4Y2	Q9t4y2 asterina pe
16	16	51.6	8	8 Q94VJ4	Q94vj4 varanus ben

17	16	51.6	8	8 Q94V91	Q94v91 varanus tim
18	16	51.6	8	8 Q94VE4	Q94ve4 varanus mel
19	16	51.6	8	8 Q94VF9	Q94vf9 varanus ind
20	16	51.6	9	8 Q9T688	Q9t688 gecko gecko
21	16	51.6	9	8 Q94VH4	Q94vh4 varanus gla
22	16	51.6	9	8 Q94VD8	Q94vd8 varanus nil
23	16	51.6	9	8 Q94VI8	Q94vi8 varanus ere
24	16	51.6	9	8 Q94VJ1	Q94vj1 varanus dor
25	16	51.6	9	8 Q8WGE6	Q8wge6 procambarus
26	16	51.6	9	8 Q94VE1	Q94ve1 varanus mer
27	16	51.6	10	2 Q93T35	Q93t35 acinetobact
28	16	51.6	10	8 Q9T8P3	Q9t8p3 liolaemus a
29	16	51.6	10	8 Q9B4W1	Q9b4w1 triturus vu
30	16	51.6	10	8 Q9T8K7	Q9t8k7 liolaemus m
31	16	51.6	10	8 Q9T8N1	Q9t8n1 liolaemus p
32	16	51.6	10	8 Q79903	Q79903 oplurus cuv
33	16	51.6	10	8 Q8WDH0	Q8wdh0 anolis limi
34	16	51.6	10	8 Q8W969	Q8w969 anolis orto
35	16	51.6	10	8 Q8WDH8	Q8wdh8 anolis mest
36	16	51.6	10	8 Q79924	Q79924 elgaria pan
37	16	51.6	10	8 Q9T8T6	Q9t8t6 liolaemus m
38	16	51.6	10	8 Q9T8L3	Q9t8l3 liolaemus l
39	16	51.6	10	8 P92616	P92616 aspidosceli
40	16	51.6	10	8 Q9T8G8	Q9t8g8 liolaemus c
41	16	51.6	10	8 Q9B4X0	Q9b4x0 notophthalm
42	16	51.6	10	8 Q8SHI3	Q8shi3 chamealeo c
43	16	51.6	10	8 Q958K9	Q958k9 rana boyllii
44	16	51.6	10	8 Q9TFU9	Q9tfu9 teratoscinc
45	16	51.6	10	8 Q9T8X7	Q9t8x7 phymaturus

ALIGNMENTS

RESULT 1  
Q9R5M1 ID Q9R5M1 PRELIMINARY; PRT; 9 AA.  
AC Q9R5M1;  
DT 01-MAY-2000 (TrEMBLrel. 13, Created)  
DT 01-MAY-2000 (TrEMBLrel. 13, Last sequence update)  
DT 01-JUN-2002 (TrEMBLrel. 21, Last annotation update)  
DE 66 kDa cell surface adhesin for heparan sulfate (Fragment).  
OS Staphylococcus aureus.  
OC Bacteria; Firmicutes; Bacillales; Staphylococcus.  
OX NCBI\_TaxID=1280;  
RN [1]  
RP SEQUENCE.  
RX MEDLINE=92176005; PubMed=1541563;  
RA Liang O.D., Ascencio F., Fransson L.A., Wadstrom T.;  
RT "Binding of heparan sulfate to Staphylococcus aureus.";  
RL Infect. Immun. 60:899-906(1992).  
FT NON\_TER 1  
FT NON\_TER 9  
SQ SEQUENCE 9 AA; 990 MW; 2289DDD7337861B3 CRC64;

Query Match 67.7%; Score 21; DB 2; Length 9;  
Best Local Similarity 50.0%; Pred. No. 8.3e+05;  
Matches 2; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

Qy 1 WXXW 4  
Db 2 WTGW 5

RESULT 2  
Q38366 ID Q38366 PRELIMINARY; PRT; 9 AA.  
AC Q38366;  
DT 01-NOV-1996 (TrEMBLrel. 01, Created)  
DT 01-NOV-1996 (TrEMBLrel. 01, Last sequence update)  
DT 01-DEC-2001 (TrEMBLrel. 19, Last annotation update)  
DE E gene product (Fragment).  
OS Bacteriophage phi-X174.



OC Viruses; ssDNA viruses; Microviridae; Microvirus.  
OX NCBI\_TaxID=10847;  
RN [1]  
RP SEQUENCE FROM N.A.  
RX MEDLINE=88118956; PubMed=2963134;  
RA Buckley K.J., Hayashi M.;  
RT "Role of premature translational termination in the regulation of  
RT expression of the phiX174 lysis gene.";  
RL J. Mol. Biol. 198:599-607(1987).  
DR EMBL; X07809; CAA30668.1; -.  
FT NON\_TER 9  
SQ SEQUENCE 9 AA; 1207 MW; C093B37731B36412 CRC64;  
  
Query Match 67.7%; Score 21; DB 9; Length 9;  
Best Local Similarity 50.0%; Pred. No. 8.3e+05;  
Matches 2; Conservative 0; Mismatches 2; Indels 0; Gaps 0;  
  
QY 1 WXXW 4  
Db 4 WTLW 7  
  
RESULT 3  
Q8SHFO  
ID Q8SHFO PRELIMINARY; PRT; 9 AA.  
AC Q8SHFO;  
DT 01-JUN-2002 (TrEMBLrel. 21, Created)  
DT 01-JUN-2002 (TrEMBLrel. 21, Last sequence update)  
DT 01-JUN-2002 (TrEMBLrel. 21, Last annotation update)  
DE Cytochrome c oxidase subunit I (Fragment).  
GN COI.  
OS Chamaeleo namaquensis.  
OG Mitochondrion.  
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
OC Lepidosauria; Squamata; Iguania; Acrodonta; Chamaeleonidae; Chamaeleo.  
OX NCBI\_TaxID=179917;  
RN [1]  
RP SEQUENCE FROM N.A.  
RA Townsend T.M., Larson A.L.;  
RT "Molecular Phylogenetics and Mitochondrial Genomic Evolution in the  
RT Chamaeleonidae (Reptilia, Squamata).";  
RL Submitted (NOV-2001) to the EMBL/GenBank/DBJ databases.  
DR EMBL; AF48757; AAL90553.1; -.  
KW Mitochondrion.  
FT NON\_TER 9  
SQ SEQUENCE 9 AA; 1205 MW; 358CB72733640733 CRC64;  
  
Query Match 64.5%; Score 20; DB 8; Length 9;  
Best Local Similarity 50.0%; Pred. No. 8.3e+05;  
Matches 2; Conservative 0; Mismatches 2; Indels 0; Gaps 0;  
  
QY 1 WXXW 4  
Db 2 WLRW 5  
  
RESULT 4  
O77919  
ID O77919 PRELIMINARY; PRT; 12 AA.  
AC O77919;  
DT 01-NOV-1998 (TrEMBLrel. 08, Created)  
DT 01-NOV-1998 (TrEMBLrel. 08, Last sequence update)  
DT 01-DEC-2001 (TrEMBLrel. 19, Last annotation update)  
DE MHC class II B locus 4 (Fragment).  
OS Pseudotropheus sp. 'Pseudotropheus tropheops complex'.  
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
OC Actinopterygii; Neopterygii; Teleostei; Euteleostei; Neoteleostei;  
OC Acanthomorpha; Acanthopterygii; Percomorpha; Perciformes; Labroidae;  
OC Cichlidae; Pseudotropheus.  
OX NCBI\_TaxID=51796;  
RN [1]  
RP SEQUENCE FROM N.A.  
RX MEDLINE=98315113; PubMed=9649539;

RA Malaga-Trillo E., Zaleska-Rutczynska Z., McAndrew B., Vincek V.,  
RA Figueroa F., Sultmann H., Klein J.;  
RT "Linkage relationships and haplotype polymorphism among cichlid mhc  
RT class II B loci.";  
RL Genetics 149:1527-1537(1998).  
DR EMBL; AF050032; AAC41371.1; -.  
FT NON\_TER 1  
FT NON\_TER 12  
SQ SEQUENCE 12 AA; 1529 MW; 6C2ABFACD5A5B734 CRC64;  
  
Query Match 64.5%; Score 20; DB 7; Length 12;  
Best Local Similarity 50.0%; Pred. No. 1.7e+03;  
Matches 2; Conservative 0; Mismatches 2; Indels 0; Gaps 0;  
  
QY 1 WXXW 4  
Db 1 WDFW 4  
  
RESULT 5  
Q16406  
ID Q16406 PRELIMINARY; PRT; 13 AA.  
AC Q16406;  
DT 01-NOV-1996 (TrEMBLrel. 01, Created)  
DT 01-NOV-1996 (TrEMBLrel. 01, Last sequence update)  
DT 01-MAY-1999 (TrEMBLrel. 10, Last annotation update)  
DE GHRH-R protein (Fragment).  
GN GHRH-R.  
OS Homo sapiens (Human).  
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
OC Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.  
OX NCBI\_TaxID=9606;  
RN [1]  
RP SEQUENCE FROM N.A.  
RX MEDLINE=96001284; PubMed=7559877;  
RA Hashimoto K., Koga M., Motomura T., Kasayama S., Kouhara H.,  
RA Ohnishi T., Arita N., Hayakawa T., Sato B., Kishimoto T.;  
RT "Identification of alternatively spliced messenger ribonucleic acid  
RT encoding truncated growth hormone-releasing hormone receptor in human  
RT pituitary adenomas.";  
RL J. Clin. Endocrinol. Metab. 80:2933-2939(1995).  
DR EMBL; S79912; AAD14318.1; -.  
FT NON\_TER 1  
SQ SEQUENCE 13 AA; 1612 MW; CB19D7D255D66362 CRC64;  
  
Query Match 64.5%; Score 20; DB 4; Length 13;  
Best Local Similarity 50.0%; Pred. No. 1.8e+03;  
Matches 2; Conservative 0; Mismatches 2; Indels 0; Gaps 0;  
  
QY 1 WXXW 4  
Db 7 WGYW 10  
  
RESULT 6  
Q53580  
ID Q53580 PRELIMINARY; PRT; 15 AA.  
AC Q53580;  
DT 01-NOV-1996 (TrEMBLrel. 01, Created)  
DT 01-NOV-1996 (TrEMBLrel. 01, Last sequence update)  
DT 01-DEC-2001 (TrEMBLrel. 19, Last annotation update)  
DE Light-harvesting complex I alpha polypeptide (Fragment).  
GN PUFA.  
OS Rhodobacter capsulatus (Rhodospseudomonas capsulata).  
OC Bacteria; Proteobacteria; Alphaproteobacteria; Rhodobacterales;  
OC Rhodobacteraceae; Rhodobacter.  
OX NCBI\_TaxID=1061;  
RN [1]  
RP SEQUENCE FROM N.A.  
RX MEDLINE=92234963; PubMed=1569029;  
RA Richter P., Brand M., Drews G.;  
RT "Characterization of LHI- and LHI+ Rhodobacter capsulatus pufa  
RT mutants.";

RL J. Bacteriol. 174:3030-3041(1992).  
DR EMBL; S97552; AAC60406.1; -.  
FT NON TER 15  
SQ SEQUENCE 15 AA; 2054 MW; 3561FE413591D31A CRC64;

Query Match 64.5%; Score 20; DB 2; Length 15;  
Best Local Similarity 50.0%; Pred. No. 2e+03;  
Matches 2; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

QY 1 WXXW 4  
Db 8 WKIW 11

RESULT 7  
Q94VC1 PRELIMINARY; PRT; 8 AA.  
AC Q94VC1;  
DT 01-DEC-2001 (TrEMBLrel. 19, Created)  
DT 01-DEC-2001 (TrEMBLrel. 19, Last sequence update)  
DT 01-DEC-2001 (TrEMBLrel. 19, Last annotation update)  
DE Cytochrome c oxidase subunit I (Fragment).  
GN COI.  
OS Varanus ruidicollis.  
OG Mitochondrion.  
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
OC Lepidosauria; Squamata; Scleroglossa; Anguimorpha; Varanidae; Varanus.  
OX NCBI\_TaxID=169851;  
RN [1]  
RP SEQUENCE FROM N.A.  
RA Ast J.C.;  
RT "Mitochondrial DNA evidence and evolution in Varanoidea (Squamata).";  
RL Cladistics 17:0-0(2001).  
DR EMBL; AF407521; AAL10116.1; -.  
KW Mitochondrion.  
FT NON TER 8  
SQ SEQUENCE 8 AA; 1053 MW; FE2729D5A36411A6 CRC64;

Query Match 54.8%; Score 17; DB 8; Length 8;  
Best Local Similarity 66.7%; Pred. No. 8.3e+05;  
Matches 2; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

QY 4 WXF 6  
Db 4 WSP 6

RESULT 8  
Q94V77 PRELIMINARY; PRT; 11 AA.  
AC Q94V77;  
DT 01-DEC-2001 (TrEMBLrel. 19, Created)  
DT 01-DEC-2001 (TrEMBLrel. 19, Last sequence update)  
DT 01-DEC-2001 (TrEMBLrel. 19, Last annotation update)  
DE Cytochrome c oxidase subunit I (Fragment).  
GN COI.  
OS Heloderma suspectum (Gila monster).  
OG Mitochondrion.  
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
OC Lepidosauria; Squamata; Scleroglossa; Anguimorpha; Helodermatidae;  
OC Heloderma.  
OX NCBI\_TaxID=8554;  
RN [1]  
RP SEQUENCE FROM N.A.  
RA Ast J.C.;  
RT "Mitochondrial DNA evidence and evolution in Varanoidea (Squamata).";  
RL Cladistics 17:0-0(2001).  
DR EMBL; AF407540; AAL10172.1; -.  
KW Mitochondrion.  
FT NON TER 11  
SQ SEQUENCE 11 AA; 1396 MW; 8E3A6DE0D5A36411 CRC64;

Query Match 54.8%; Score 17; DB 8; Length 11;

Best Local Similarity 66.7%; Pred. No. 4.8e+03;  
Matches 2; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

QY 4 WXF 6  
Db 6 WSP 8

RESULT 9  
Q9UDC6 PRELIMINARY; PRT; 13 AA.  
ID Q9UDC6  
AC Q9UDC6;  
DT 01-MAY-2000 (TrEMBLrel. 13, Created)  
DT 01-MAY-2000 (TrEMBLrel. 13, Last sequence update)  
DT 01-JUN-2002 (TrEMBLrel. 21, Last annotation update)  
DE ENDOTHELIUM-derived RELATING factor, nitric oxide synthase (Fragment).  
DE Homo sapiens (Human).  
OS Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
OC Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.  
OX NCBI\_TaxID=9606;  
RN [1]  
RP SEQUENCE FROM N.A.  
RX MEDLINE=93054573; PubMed=1385404;  
RA Janssens S.P., Simouchi A., Quertermous T., Bloch D.B., Bloch K.D.;  
RT "Cloning and expression of a cDNA encoding human endothelium-derived relating factor/nitric oxide synthase.";  
RL J. Biol. Chem. 267:22694-22694(1992).  
FT NON TER 1  
FT NON TER 13  
SQ SEQUENCE 13 AA; 1390 MW; 3231B6DFEC7EB867 CRC64;

Query Match 54.8%; Score 17; DB 4; Length 13;  
Best Local Similarity 66.7%; Pred. No. 5.5e+03;  
Matches 2; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

QY 4 WXF 6  
Db 1 WAF 3

RESULT 10  
Q9SAP8 PRELIMINARY; PRT; 14 AA.  
ID Q9SAP8  
AC Q9SAP8;  
DT 01-MAY-2000 (TrEMBLrel. 13, Created)  
DT 01-MAY-2000 (TrEMBLrel. 13, Last sequence update)  
DT 01-MAY-2000 (TrEMBLrel. 13, Last annotation update)  
DE LHCPII (14AA) (Fragment).  
OS Pisum sativum (Garden pea).  
OC Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;  
OC Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; Rosidae;  
OC eurosids I; Fabales; Fabaceae; Papilionoideae; Viciae; Pisum.  
OX NCBI\_TaxID=3888;  
RN [1]  
RP SEQUENCE FROM N.A.  
RC STRAIN=var. Alaska;  
RA Dobres M.S., Abler M.L., Thompson W.F.;  
RT "Sequence of the 3' untranslated region of a pea.";  
RL Nucleic Acids Res. 0:0-0(1988).  
DR EMBL; X06822; CAA29970.1; -.  
FT NON TER 1  
SQ SEQUENCE 14 AA; 1537 MW; D55621E9906EA7AD CRC64;

Query Match 54.8%; Score 17; DB 10; Length 14;  
Best Local Similarity 66.7%; Pred. No. 5.8e+03;  
Matches 2; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

QY 4 WXF 6  
Db 4 WAF 6

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RESULT 11
Q94VF6          PRELIMINARY;      PRT;      8 AA.
AC  Q94VF6;
DT  01-DEC-2001 (TrEMBLrel. 19, Created)
DT  01-DEC-2001 (TrEMBLrel. 19, Last sequence update)
DT  01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
DE  Cytochrome c oxidase subunit I (Fragment).
GN  COI.
OS  Varanus jobiensis.
OG  Mitochondrion.
OC  Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC  Lepidosauria; Squamata; Scleroglossa; Anguimorpha; Varanidae; Varanus.
OX  NCBI_TaxID=169843;
RN  [1]
RP  SEQUENCE FROM N.A.
RA  Ast J.C.;
RT  "Mitochondrial DNA evidence and evolution in Varanoidea (Squamata).";
RL  Cladistics 17:0-0(2001).
DR  EMBL; AF407507; AAL10075.1; -.
KW  Mitochondrion.
FT  NON_TER      8
SQ  SEQUENCE      8 AA; 1144 MW; EFD729DB436411A6 CRC64;

  Query Match      51.6%; Score 16; DB 8; Length 8;
  Best Local Similarity 66.7%; Pred. No. 8.3e+05;
  Matches 2; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

Qy      4 WXF 6
Db      4 WYF 6

RESULT 12
Q8WGD7          PRELIMINARY;      PRT;      8 AA.
AC  Q8WGD7;
DT  01-MAR-2002 (TrEMBLrel. 20, Created)
DT  01-MAR-2002 (TrEMBLrel. 20, Last sequence update)
DT  01-MAR-2002 (TrEMBLrel. 20, Last annotation update)
DE  Cytochrome oxidase subunit 1 (Fragment).
OS  Lomis hirta.
OG  Mitochondrion.
OC  Eukaryota; Metazoa; Arthropoda; Crustacea; Malacostraca;
OC  Eumalacostraca; Eucarida; Decapoda; Pleocyemata; Anomura; Lomoidea;
OC  Lomidae; Lomis.
OX  NCBI_TaxID=177234;
RN  [1]
RP  SEQUENCE FROM N.A.
RA  Morrison C.L., Harvey A.W., Lavery S., Tieu K., Huang Y.,
RA  Cunningham C.W.;
RT  "Mitochondrial gene rearrangements support a hypothesis of parallel
RT  evolution to the crab-like form.";
RL  Submitted (OCT-2001) to the EMBL/GenBank/DBJ databases.
DR  EMBL; AF436035; AAL31611.1; -.
KW  Mitochondrion.
FT  NON_TER      1
FT  NON_TER      8
SQ  SEQUENCE      8 AA; 1038 MW; C5B5B9C733640321 CRC64;

  Query Match      51.6%; Score 16; DB 8; Length 8;
  Best Local Similarity 66.7%; Pred. No. 8.3e+05;
  Matches 2; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

Qy      4 WXF 6
Db      4 WLF 6

RESULT 13
Q94V88          PRELIMINARY;      PRT;      8 AA.
AC  Q94V88;

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DT  01-DEC-2001 (TrEMBLrel. 19, Created)
DT  01-DEC-2001 (TrEMBLrel. 19, Last sequence update)
DT  01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
DE  Cytochrome c oxidase subunit I (Fragment).
GN  COI.
OS  Varanus tristis.
OG  Mitochondrion.
OC  Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC  Lepidosauria; Squamata; Scleroglossa; Anguimorpha; Varanidae; Varanus.
OX  NCBI_TaxID=62052;
RN  [1]
RP  SEQUENCE FROM N.A.
RA  Ast J.C.;
RT  "Mitochondrial DNA evidence and evolution in Varanoidea (Squamata).";
RL  Cladistics 17:0-0(2001).
DR  EMBL; AF407533; AAL10151.1; -.
KW  Mitochondrion.
FT  NON_TER      8
SQ  SEQUENCE      8 AA; 1041 MW; E8B5B9C7336411A6 CRC64;

  Query Match      51.6%; Score 16; DB 8; Length 8;
  Best Local Similarity 66.7%; Pred. No. 8.3e+05;
  Matches 2; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

Qy      4 WXF 6
Db      4 WLF 6

RESULT 14
Q9TD02          PRELIMINARY;      PRT;      8 AA.
AC  Q9TD02;
DT  01-MAY-2000 (TrEMBLrel. 13, Created)
DT  01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
DT  01-MAY-2000 (TrEMBLrel. 13, Last annotation update)
DE  Cytochrome c oxidase subunit I (Fragment).
OS  Terranatos dolichopterus.
OG  Mitochondrion.
OC  Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC  Actinopterygii; Neopterygii; Teleostei; Euteleostei; Neoteleostei;
OC  Acanthomorpha; Acanthopterygii; Percomorpha; Atherinomorpha;
OC  Cyprinodontiformes; Aplocheilidae; Rivulinae; Terranatos.
OX  NCBI_TaxID=61836;
RN  [1]
RP  SEQUENCE FROM N.A.
RA  Hrbek T., Larson A.;
RT  "The evolution of diapause in the killifish family Rivulidae
RT  (Atherinomorpha, Cyprinodontiformes): A molecular phylogenetic and
RT  biogeographic perspective.";
RL  Evolution 53:1200-1216(1999).
DR  EMBL; AF092421; AAF03041.1; -.
KW  Mitochondrion.
FT  NON_TER      8
SQ  SEQUENCE      8 AA; 1084 MW; F0C9D3640DD44056 CRC64;

  Query Match      51.6%; Score 16; DB 8; Length 8;
  Best Local Similarity 66.7%; Pred. No. 8.3e+05;
  Matches 2; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

Qy      4 WXF 6
Db      6 WFF 8

RESULT 15
Q9T4Y2          PRELIMINARY;      PRT;      8 AA.
AC  Q9T4Y2;
DT  01-MAY-2000 (TrEMBLrel. 13, Created)
DT  01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
DT  01-MAY-2000 (TrEMBLrel. 13, Last annotation update)
DE  COI gene product (Fragment).

```

OS Asterina pectinifera (Starfish).  
OG Mitochondrion.  
OC Eukaryota; Metazoa; Echinodermata; Eleutherozoa; Asterozoa;  
OC Asteroidea; Valvatacea; Valvatida; Asterinidae; Asterina.  
OX NCBI\_TaxID=7594;  
RN [1]  
RP SEQUENCE FROM N.A.  
RX MEDLINE=89354669; PubMed=2766382;  
RA Jacobs H.T.; Asakawa S.; Araki T.; Miura K.; Smith M.J.; Watanabe K.;  
RT "Conserved tRNA gene cluster in starfish mitochondrial DNA.";  
RL Curr. Genet. 15:193-206(1989).  
DR EMBL; X16886; CAA34767.1; -.  
KW Mitochondrion.  
FT NON TER 8  
SQ SEQUENCE 8 AA; 1114 MW; F0C9D36415B736D6 CRC64;

Query Match 51.6%; Score 16; DB 8; Length 8;  
Best Local Similarity 66.7%; Pred. No. 8.3e+05;  
Matches 2; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

QY 4 WXF 6  
| |  
Db 6 WFF 8

Search completed: December 3, 2003, 11:53:24  
Job time : 27.3333 secs

GenCore version 5.1.6  
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OM protein - protein search, using sw model

Run on: December 3, 2003, 11:44:55 ; Search time 7.33333 Seconds  
(without alignments)  
38.476 Million cell updates/sec

Title: US-09-912-414-9  
Perfect score: 31  
Sequence: 1 WXXWXF 6

Scoring table: BLOSUM62  
Gapop 10.0 , Gapext 0.5

Searched: 127863 seqs, 47026705 residues

Total number of hits satisfying chosen parameters: 795

Minimum DB seq length: 0  
Maximum DB seq length: 15

Post-processing: Minimum Match 0%  
Maximum Match 100%  
Listing first 45 summaries

Database : SwissProt\_41:\*

Pred. No. is the number of results predicted by chance to have a  
score greater than or equal to the score of the result being printed,  
and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	DB ID	Description
1	21	67.7	10	1 LABA JATMU	P13270 jatropha mu
2	19	61.3	9	1 COW CONVE	P83047 conus ventr
3	14	45.2	9	1 LITR PHYRO	P08946 phyllomedus
4	14	45.2	10	1 GON1 ALLMI	P37041 alligator m
5	14	45.2	10	1 GON3 ONCKE	P20367 oncorhynchu
6	14	45.2	11	1 RANC RANPI	P08951 rana pipien
7	13	41.9	8	1 RT34 BOVIN	P82929 bos taurus
8	13	41.9	9	1 LITO LITAU	P08945 litoria aur
9	13	41.9	10	1 HTF TABAT	P14596 tabanus atr
10	13	41.9	12	1 UR2A CATCO	P04558 catostomus
11	13	41.9	12	1 UR2B CATCO	P04559 catostomus
12	13	41.9	12	1 UR2B CYPCA	P04561 cyprinus ca
13	13	41.9	12	1 UR2 GILMI	P01147 gillichthys
14	13	41.9	12	1 UR2 POLSP	P81022 polyodon sp
15	13	41.9	12	1 UR2 SCYCA	P35490 scylliorhinu
16	13	41.9	13	1 BOML PSEGU	P42991 pseudophryn
17	12	38.7	6	1 LOK1 LOCFI	P41491 locusta mig
18	12	38.7	8	1 LCK2 LEUMA	P21141 leucophaea
19	12	38.7	8	1 LCK5 LEUMA	P19987 leucophaea
20	12	38.7	8	1 LCK7 LEUMA	P19989 leucophaea
21	12	38.7	10	1 AEGL AGRAE	P83465 agroclybe ae
22	12	38.7	10	1 CA12 LITCI	P82086 litoria cit
23	12	38.7	10	1 CAER LITXA	P56264 litoria xan
24	12	38.7	10	1 GON1 CHEPR	P80677 chelyosoma
25	12	38.7	13	1 YPNP PHOLU	P41122 photorhabdu
26	12	38.7	15	1 RM12 YEAST	P36522 saccharomyc
27	11	35.5	4	1 OCP3 OCTMI	P58649 octopus min
28	11	35.5	5	1 BPP7 BOTIN	P30425 bothrops in
29	11	35.5	5	1 UF01 MOUSE	P38639 mus musculu
30	11	35.5	6	1 EI01 LITRU	P82096 litoria rub
31	11	35.5	7	1 BRHP CONIM	P58803 conus imper
32	11	35.5	7	1 TPFY PACDA	P83455 pachymedusa
33	11	35.5	7	1 TY51 LITRU	P82065 litoria rub

34	11	35.5	7	1 WWA1 ACHFUF	P35919 achatina fu
35	11	35.5	7	1 WWA2 ACHFUF	P35920 achatina fu
36	11	35.5	7	1 WWA3 ACHFUF	P35921 achatina fu
37	11	35.5	8	1 ACI THUAL	P18691 thunnus alb
38	11	35.5	8	1 AKHG GRVBI	P14086 gryllus bim
39	11	35.5	8	1 AKH LIAU	P25418 libellula a
40	11	35.5	8	1 AKH MELML	P25423 melolontha
41	11	35.5	8	1 AKH TABAT	P14595 tabanus atr
42	11	35.5	8	1 CCKN MACPU	P30369 macropus eu
43	11	35.5	8	1 COW2 CONPU	P58785 conus purpu
44	11	35.5	8	1 HTF1 PERAM	P04548 periplaneta
45	11	35.5	8	1 HTF2 PERAM	P04549 periplaneta

ALIGNMENTS

RESULT 1  
LABA JATMU  
ID LABA JATMU STANDARD; PRT; 10 AA.  
AC P13270;  
DT 01-JAN-1990 (Rel. 13, Created)  
DT 01-JAN-1990 (Rel. 13, Last sequence update)  
DT 28-FEB-2003 (Rel. 41, Last annotation update)  
DE Labaditin.  
OS Jatropha multifida (Physic nut).  
OC Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;  
OC Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; Rosidae;  
OC eurosids I; Malpighiales; Euphorbiaceae; Jatropha.  
OX NCBI\_TaxID=3996;  
RN [1]  
RP SEQUENCE.  
RC TISSUE=Latex;  
RA Kosasi S., van der Sluis W.G., Boelens R., T'Hart L.A., Labadie R.P.;  
RT "Labaditin, a novel cyclic decapeptide from the latex of Jatropha  
multifida L. (Euphorbiaceae). Isolation and sequence determination  
by means of two-dimensional NMR.";  
RL FEBS Lett. 256:91-96(1989).  
CC -!- FUNCTION: LABADITIN IS AN ACTIVE PEPTIDE WHICH INHIBITS THE  
CLASSICAL PATHWAY OF COMPLEMENT ACTIVATION IN VITRO. ACTIVITY  
SEEMS TO BE BASED ON AN INTERACTION WITH C1.  
CC -!- PTM: This is a cyclic peptide.  
CC -!- DISEASE: LATEX OF THIS PLANT IS USED IN FOLKLORIC MEDICINE FOR  
TREATMENT OF INFECTED WOUNDS, SKINS INFECTIONS AND SCABIES.  
SQ SEQUENCE 10 AA; 1089 MW; D98AAD6362D1B362 CRC64;  
Query Match 67.7%; Score 21; DB 1; Length 10;  
Best Local Similarity 50.0%; Pred.No. 1.7e+02;  
Matches 2; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

QY	1	WXXW	4
Db	4	WTVW	7

RESULT 2  
COW CONVE  
ID COW CONVE STANDARD; PRT; 9 AA.  
AC P83047;  
DT 16-OCT-2001 (Rel. 40, Created)  
DT 16-OCT-2001 (Rel. 40, Last sequence update)  
DT 28-FEB-2003 (Rel. 41, Last annotation update)  
DE Contryphan-Vn.  
OS Conus ventricosus (Mediterranean cone).  
OC Eukaryota; Metazoa; Mollusca; Gastropoda; Orthogastropoda;  
OC Apogastropoda; Caenogastropoda; Sorbeoconcha; Hypsogastropoda;  
OC Neogastropoda; Conoidea; Conidae; Conus.  
OX NCBI\_TaxID=117992;  
RN [1]  
RP SEQUENCE, SYNTHESIS, AND MASS SPECTROMETRY.  
RC TISSUE=Venom;  
RX MEDLINE=21547785; PubMed=11688995;  
RA Massilia G.R., Schinina M.E., Ascenzi P., Polticelli F.;



RT "Contryphan-Vn: a novel peptide from the venom of the Mediterranean  
RT snail *Conus ventricosus*,"  
RL Biochem. Biophys. Res. Commun. 288:908-913(2001).  
CC -!- SUBCELLULAR LOCATION: Secreted.  
CC -!- TISSUE SPECIFICITY: Expressed by the venom duct.  
CC -!- MASS SPECTROMETRY: MW=1088.6; METHOD=MALDI.  
CC -!- SIMILARITY: BELONGS TO THE CONTRYPHAN FAMILY.  
KW Toxin; Amidation; D-amino acid.  
FT DISULFID 3 9  
FT MOD\_RES 5 5 D-TRYPTOPHAN.  
FT MOD\_RES 9 9 AMIDATION.  
SQ SEQUENCE 9 AA; 1091 MW; 8D38676323676EBA CRC64;  
  
Query Match 61.3%; Score 19; DB 1; Length 9;  
Best Local Similarity 50.0%; Pred. No. 1.3e+05;  
Matches 2; Conservative 0; Mismatches 2; Indels 0; Gaps 0;  
  
QY 1 WXXW 4  
Db 5 WKPW 8  
  
RESULT 3  
LITR\_PHYRO STANDARD; PRT; 9 AA.  
AC P08946;  
DT 01-NOV-1988 (Rel. 09, Created)  
DT 01-FEB-1994 (Rel. 28, Last sequence update)  
DT 15-SEP-2003 (Rel. 42, Last annotation update)  
DE Rhodelalitorin.  
OS Phyllomedusa rohdei (Rohde's leaf frog).  
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
OC Amphibia; Batrachia; Anura; Neobatrachia; Bufonoidea; Hylidae;  
OC Phyllomedusinae; Phyllomedusa.  
OX NCBI\_TaxID=8394;  
RN [1]  
RP SEQUENCE.  
RC TISSUE=Skin secretion;  
RX MEDLINE=85127560; PubMed=3838283;  
RA Barra D., Erspamer G.F., Simmaco M., Bossa F., Melchiorri P.,  
RA Erspamer V.;  
RT "Rohdei-litorin: a new peptide from the skin of Phyllomedusa rohdei,";  
RL FEBS Lett. 182:53-56(1985).  
CC -!- SUBCELLULAR LOCATION: Secreted.  
CC -!- TISSUE SPECIFICITY: Skin.  
CC -!- SIMILARITY: BELONGS TO THE BOMBESIN/NEUROMEDIN B/RANATENSIN  
CC FAMILY.  
DR PIR; S07241; S07241.  
DR InterPro; IPR000874; Bombesin.  
DR Pfam; PF02044; Bombesin; 1.  
DR PROSITE; PS00257; BOMBESIN; 1.  
KW Amphibian defense peptide; Bombesin family; Amidation;  
KW Pyrrolidone carboxylic acid.  
FT MOD\_RES 1 1 PYRROLIDONE CARBOXYLIC ACID.  
FT MOD\_RES 9 9 AMIDATION.  
SQ SEQUENCE 9 AA; 1090 MW; 4ECCC1E861ADC377 CRC64;  
  
Query Match 45.2%; Score 14; DB 1; Length 9;  
Best Local Similarity 33.3%; Pred. No. 1.3e+05;  
Matches 2; Conservative 0; Mismatches 4; Indels 0; Gaps 0;  
  
QY 1 WXXW 6  
Db 3 WATG 8  
  
RESULT 4  
GON1\_ALLMI STANDARD; PRT; 10 AA.  
AC P37041; P20407;  
DT 01-FEB-1991 (Rel. 17, Created)  
DT 01-FEB-1991 (Rel. 17, Last sequence update)  
DT 28-FEB-2003 (Rel. 41, Last annotation update)

DE Gonadoliberin I (Gonadotropin-releasing hormone I) (GnRH-I) (LH-RH I)  
DE (Luliberin I).  
OS Alligator mississippiensis (American alligator).  
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
OC Archosauria; Crocodylidae; Alligatorinae; Alligator.  
OX NCBI\_TaxID=8496;  
RN [1]  
RP SEQUENCE.  
RC TISSUE=Brain;  
RX MEDLINE=91352338; PubMed=1882082;  
RA Lovejoy D.A., Fischer W.H., Parker D.B., McRory J.E., Park M.,  
RA Lance V., Swanson P., Rivier J.E., Sherwood N.M.;  
RT "Primary structure of two forms of gonadotropin-releasing hormone  
from brains of the American alligator (Alligator mississippiensis).";  
RL Regul. Pept. 33:105-116(1991).  
CC -!- FUNCTION: Stimulates the secretion of gonadotropins.  
CC -!- SUBCELLULAR LOCATION: Secreted.  
CC -!- SIMILARITY: Belongs to the GnRH family.  
DR PIR; A60066; RHAQ1.  
DR InterPro; IPR002012; GnRH.  
DR Pfam; PF00446; GnRH; 1.  
DR PROSITE; PS00473; GnRH; 1.  
KW Hormone; Amidation; Hypothalamus; Pyrrolidone carboxylic acid.  
FT MOD\_RES 1 1 PYRROLIDONE CARBOXYLIC ACID.  
FT MOD\_RES 10 10 AMIDATION.  
SQ SEQUENCE 10 AA; 1172 MW; 284B23D7286B45A3 CRC64;  
  
Query Match 45.2%; Score 14; DB 1; Length 10;  
Best Local Similarity 33.3%; Pred. No. 2.4e+03;  
Matches 1; Conservative 1; Mismatches 1; Indels 0; Gaps 0;  
  
QY 4 WXF 6  
Db 3 WSY 5  
  
RESULT 5  
GON3\_ONCKE STANDARD; PRT; 10 AA.  
AC P20367; P81751;  
DT 01-FEB-1991 (Rel. 17, Created)  
DT 01-FEB-1991 (Rel. 17, Last sequence update)  
DT 28-FEB-2003 (Rel. 41, Last annotation update)  
DE Gonadoliberin III (Gonadotropin-releasing hormone III) (GnRH-III) (LH-  
RH III) (Luliberin III).  
GN GnRH3.  
OS Oncorhynchus keta (Chum salmon), and  
OS Clupea pallasii (Pacific herring).  
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
OC Actinopterygii; Neopterygii; Teleostei; Euteleostei;  
OC Protacanthopterygii; Salmoniformes; Salmonidae; Oncorhynchus.  
OX NCBI\_TaxID=8018, 30724;  
RN [1]  
RP SEQUENCE.  
RC SPECIES=O.keta;  
RX MEDLINE=83195140; PubMed=6341999;  
RA Sherwood N., Eiden L., Brownstein M., Spiess J., Rivier J., Vale W.;  
RT "Characterization of a teleost gonadotropin-releasing hormone.";  
RL Proc. Natl. Acad. Sci. U.S.A. 80:2794-2798(1983).  
RN [2]  
RP SEQUENCE, AND FUNCTION.  
RC SPECIES=C.pallasii; TISSUE=Brain, and Pituitary;  
RX MEDLINE=20114351; PubMed=10650929;  
RA Carlsfeld J., Powell J.F.F., Park M., Fischer W.H., Craig A.G.,  
RA Chang J.P., Rivier J.E., Sherwood N.M.;  
RT "Primary structure and function of three gonadotropin-releasing  
hormones, including a novel form, from an ancient teleost, herring.";  
RL Endocrinology 141:505-512(2000).  
CC -!- FUNCTION: Stimulates the secretion of gonadotropins; it stimulates  
the secretion of both luteinizing and follicle-stimulating  
hormones.  
CC -!- SUBCELLULAR LOCATION: Secreted.  
CC -!- SIMILARITY: Belongs to the GnRH family.

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DR PIR; A21114; A21114.
DR InterPro; IPR002012; GnrH.
DR Pfam; PF00446; GnrH; 1.
DR PROSITE; PS00473; GNRH; 1.
KW Hormone; Amidation; Hypothalamus; Pyrrolidone carboxylic acid.
FT MOD_RES 1 1 PYRROLIDONE CARBOXYLIC ACID.
FT MOD_RES 10 10 AMIDATION.
SQ SEQUENCE 10 AA; 1230 MW; 284B3233786B45A3 CRC64;

Query Match 45.2%; Score 14; DB 1; Length 10;
Best Local Similarity 33.3%; Pred. No. 2.4e+03;
Matches 1; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

QY 4 WXF 6
Db 3 WSY 5

RESULT 6
RANC_RANPI STANDARD; PRT; 11 AA.
AC P08951;
DT 01-NOV-1988 (Rel. 09, Created)
DT 01-NOV-1988 (Rel. 09, Last sequence update)
DT 15-SEP-2003 (Rel. 42, Last annotation update)
DE Ranatensin-C.
OS Rana pipiens (Northern leopard frog).
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Amphibia; Batrachia; Anura; Neobatrachia; Ranoidea; Ranidae; Rana.
OX NCBI_TaxID=8404;
RN [1]
RP SEQUENCE.
RC TISSUE=Skin secretion;
RX MEDLINE=84131098; PubMed=6141890;
RA Nakajima T.;
RL Unpublished results, cited by:
RL Erspamer V., Erspamer G.F., Mazzanti G., Endean R.;
RL Comp. Biochem. Physiol. 77C:99-108(1984).
CC -!- SUBCELLULAR LOCATION: Secreted.
CC -!- TISSUE SPECIFICITY: Skin.
CC -!- SIMILARITY: BELONGS TO THE BOMBESIN/NEUROMEDIN B/RANATENSIN
CC FAMILY.
DR InterPro; IPR000874; Bombesin.
DR Pfam; PF02044; Bombesin; 1.
DR PROSITE; PS00257; BOMBESIN; 1.
KW Amphibian defense peptide; Bombesin family; Amidation.
FT MOD_RES 11 11 AMIDATION.
FT MOD_RES 11 11
SQ SEQUENCE 11 AA; 1304 MW; D6C9885A61ADC366 CRC64;

Query Match 45.2%; Score 14; DB 1; Length 11;
Best Local Similarity 33.3%; Pred. No. 2.6e+03;
Matches 2; Conservative 0; Mismatches 4; Indels 0; Gaps 0;

QY 1 WXXWXP 6
Db 5 WATGHF 10

RESULT 7
RT34_BOVIN STANDARD; PRT; 8 AA.
AC P82929;
DT 28-FEB-2003 (Rel. 41, Created)
DT 28-FEB-2003 (Rel. 41, Last sequence update)
DT 28-FEB-2003 (Rel. 41, Last annotation update)
DE Mitochondrial 28S ribosomal protein S34 (S34mt) (MRP-S34) (Fragment).
GN MRPS34.
OS Bos taurus (Bovine).
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Mammalia; Eutheria; Cetartiodactyla; Ruminantia; Pecora; Bovoidea;
OC Bovidae; Bovinae; Bos.
OX NCBI_TaxID=9913;
RN [1]
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RP SEQUENCE.
RC TISSUE=Liver;
RX MEDLINE=21276436; PubMed=11279123;
RA Koc E.C., Burkhardt W., Blackburn K., Moseley A., Spremulli L.L.;
RT "The small subunit of the mammalian mitochondrial ribosome:
RT identification of the full complement of ribosomal proteins present.";
RL J. Biol. Chem. 276:19363-19374(2001).
CC -!- SUBUNIT: Component of the mitochondrial ribosome small subunit
CC (28S) which comprises a 12S rRNA and about 30 distinct proteins.
CC -!- SUBCELLULAR LOCATION: Mitochondrial.
KW Ribosomal protein; Mitochondrion.
FT NON_TER 1 1
FT NON_TER 8 8
SQ SEQUENCE 8 AA; 935 MW; 9639D1A72058637D CRC64;

Query Match 41.9%; Score 13; DB 1; Length 8;
Best Local Similarity 33.3%; Pred. No. 1.3e+05;
Matches 2; Conservative 0; Mismatches 4; Indels 0; Gaps 0;

QY 1 WXXWXP 6
Db 2 WGILTF 7

RESULT 8
LITO_LITAU STANDARD; PRT; 9 AA.
AC P08945;
DT 01-NOV-1988 (Rel. 09, Created)
DT 01-FEB-1994 (Rel. 28, Last sequence update)
DT 15-SEP-2003 (Rel. 42, Last annotation update)
DE Litorin.
OS Litoria aurea (Green and golden bell frog).
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Amphibia; Batrachia; Anura; Neobatrachia; Bufonoidea; Hylidae;
OC Pelodyadinae; Litoria.
OX NCBI_TaxID=8371;
RN [1]
RP SEQUENCE.
RC TISSUE=Skin secretion;
RX MEDLINE=75187011; PubMed=1140241;
RA Anastasi A., Erspamer V., Endean R.;
RT "Aminoacid composition and sequence of litorin, a bombesin-like
RT nonapeptide from the skin of the Australian leptodactylid frog
RT Litoria aurea.";
RL Experientia 31:510-511(1975).
RN [2]
RP SEQUENCE (METHYLATED VARIANT).
RC TISSUE=Skin secretion;
RX MEDLINE=78003546; PubMed=908397;
RA Anastasi A., Montecucchi P.C., Angelucci F., Erspamer V., Endean R.;
RT "Glu(OMe)3-litorin, the second bombesin-like peptide occurring in
RT methanol extracts of the skin of the Australian frog Litoria aurea.";
RL Experientia 33:1289-1289(1977).
CC -!- SUBCELLULAR LOCATION: Secreted.
CC -!- TISSUE SPECIFICITY: Skin.
CC -!- SIMILARITY: BELONGS TO THE BOMBESIN/NEUROMEDIN B/RANATENSIN
CC FAMILY.
DR PIR; S07204; S07204.
DR InterPro; IPR000874; Bombesin.
DR Pfam; PF02044; Bombesin; 1.
DR PROSITE; PS00257; BOMBESIN; 1.
KW Amphibian defense peptide; Bombesin family; Amidation; Methylation;
KW Pyrrolidone carboxylic acid.
FT MOD_RES 1 1 PYRROLIDONE CARBOXYLIC ACID.
FT MOD_RES 2 2 METHYLATION (PARTIAL).
FT MOD_RES 9 9 AMIDATION.
FT MOD_RES 9 9
SQ SEQUENCE 9 AA; 1103 MW; D7CCC1E862CDC366 CRC64;

Query Match 41.9%; Score 13; DB 1; Length 9;
Best Local Similarity 33.3%; Pred. No. 1.3e+05;
Matches 2; Conservative 0; Mismatches 4; Indels 0; Gaps 0;
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QY 1 WXXWXF 6  
|  
Db 3 WVGWGF 8

RESULT 9  
HTF\_TABAT  
ID HTF TABAT STANDARD; PRT; 10 AA.  
AC P14596;  
DT 01-JAN-1990 (Rel. 13, Created)  
DT 01-FEB-1994 (Rel. 28, Last sequence update)  
DT 28-FEB-2003 (Rel. 41, Last annotation update)  
DE Hypertrehalosaemic factor (HOTH) (Dipteran corpora cardiaca factor II) (DCC II).  
OS Tabanus atratus (Horse fly).  
OC Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;  
OC Neoptera; Endopterygota; Diptera; Brachycera; Tabanomorpha; Tabanidae;  
OC Tabanus.  
OX NCBI\_TaxID=7207;  
RN [1]  
RP SEQUENCE.  
RC TISSUE=Corpora cardiaca;  
RX MEDLINE=90046758; PubMed=2813385;  
RA Jaffe H., Raina A.K., Riley C.T., Fraser B.A., Nachman R.J.,  
RA Vogel V.W., Zhang Y.-S., Hayes D.K.;  
RT "Primary structure of two neuropeptide hormones with adipokinetic and  
RT hypotrehalosemic activity isolated from the corpora cardiaca of horse  
RT flies (Diptera).";  
RL Proc. Natl. Acad. Sci. U.S.A. 86:8161-8164(1989).  
CC -!- FUNCTION: HYPERTREHALOSAEMIC FACTORS ARE NEUROPEPTIDES THAT  
CC ELEVATE THE LEVEL OF TREHALOSE IN THE HEMOLYMPH (TREHALOSE IS  
CC THE MAJOR CARBOHYDRATE IN THE HEMOLYMPH OF INSECTS).  
CC -!- SIMILARITY: BELONGS TO THE AKH / HRTH / RPCH FAMILY.  
DR PIR; B33995; B33995.  
DR InterPro; IPR002047; AKH.  
DR PROSITE; PS00256; AKH; 1.  
KW Neuropeptide; Amidation; Pyrrolidone carboxylic acid.  
FT MOD\_RES 1 1 PYRROLIDONE CARBOXYLIC ACID.  
FT MOD\_RES 10 10 AMIDATION.  
SQ SEQUENCE 10 AA; 1169 MW; 916036786771A9D1 CRC64;.

Query Match 41.9%; Score 13; DB 1; Length 10;  
Best Local Similarity 33.3%; Pred. No. 3.5e+03;  
Matches 1; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

QY 4 WXF 6  
|  
Db 8 WGY 10

RESULT 10  
UR2A\_CATCO  
ID UR2A CATCO STANDARD; PRT; 12 AA.  
AC P04558;  
DT 13-AUG-1987 (Rel. 05, Created)  
DT 13-AUG-1987 (Rel. 05, Last sequence update)  
DT 16-OCT-2001 (Rel. 40, Last annotation update)  
DE Urotensin IIA (U-IIA) (UIIA).  
OS Catostomus commersoni (White sucker).  
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
OC Actinopterygii; Neopterygii; Teleostei; Ostariophysi; Cypriniformes;  
OC Catostomidae; Catostomus.  
OX NCBI\_TaxID=7971;  
RN [1]  
RP SEQUENCE.  
RX MEDLINE=84041959; PubMed=6138758;  
RA McMaster D., Lederis K.;  
RT "Isolation and amino acid sequence of two urotensin II peptides from  
RT Catostomus commersoni urophyses.";  
RL Peptides 4:367-373(1983).  
CC -!- FUNCTION: UROTENSIN IS FOUND IN THE TELEOST CAUDAL NEUROSECRETORY  
CC SYSTEM. IT HAS A SUGGESTED ROLE IN OSMOREGULATION AND AS A  
CC CORTICOTROPIN-RELEASING FACTOR.

Query Match 41.9%; Score 13; DB 1; Length 12;  
Best Local Similarity 33.3%; Pred. No. 4e+03;  
Matches 1; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

QY 4 WXF 6  
|  
Db 8 WKY 10

RESULT 11  
UR2B\_CATCO  
ID UR2B CATCO STANDARD; PRT; 12 AA.  
AC P04559;  
DT 13-AUG-1987 (Rel. 05, Created)  
DT 13-AUG-1987 (Rel. 05, Last sequence update)  
DT 16-OCT-2001 (Rel. 40, Last annotation update)  
DE Urotensin IIB (U-IIB) (UIIB).  
OS Catostomus commersoni (White sucker).  
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
OC Actinopterygii; Neopterygii; Teleostei; Ostariophysi; Cypriniformes;  
OC Catostomidae; Catostomus.  
OX NCBI\_TaxID=7971;  
RN [1]  
RP SEQUENCE.  
RX MEDLINE=84041959; PubMed=6138758;  
RA McMaster D., Lederis K.;  
RT "Isolation and amino acid sequence of two urotensin II peptides from  
RT Catostomus commersoni urophyses.";  
RL Peptides 4:367-373(1983).  
CC -!- FUNCTION: UROTENSIN IS FOUND IN THE TELEOST CAUDAL NEUROSECRETORY  
CC SYSTEM. IT HAS A SUGGESTED ROLE IN OSMOREGULATION AND AS A  
CC CORTICOTROPIN-RELEASING FACTOR.

Query Match 41.9%; Score 13; DB 1; Length 12;  
Best Local Similarity 33.3%; Pred. No. 4e+03;  
Matches 1; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

QY 4 WXF 6  
|  
Db 8 WKY 10

RESULT 12  
UR2B\_CYPCA  
ID UR2B CYPCA STANDARD; PRT; 12 AA.  
AC P04561;  
DT 13-AUG-1987 (Rel. 05, Created)  
DT 13-AUG-1987 (Rel. 05, Last sequence update)  
DT 16-OCT-2001 (Rel. 40, Last annotation update)  
DE Urotensin II-beta.  
OS Cyprinus carpio (Common carp).  
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
OC Actinopterygii; Neopterygii; Teleostei; Ostariophysi; Cypriniformes;  
OC Cyprinidae; Cyprinus.  
OX NCBI\_TaxID=7962;  
RN [1]

CC -!- SIMILARITY: BELONGS TO THE UROTENSIN 2 FAMILY.  
DR PIR; JS0423; JS0423.  
DR InterPro; IPR001483; Urotensin\_II.  
DR Pfam; PF02083; Urotensin\_II; 1.  
DR PROSITE; PS00984; UROTENSIN\_II; 1.  
KW Hormone.  
FT DISULFID 6 11  
SQ SEQUENCE 12 AA; 1336 MW; 969C76DBB879CEBA CRC64;

Query Match 41.9%; Score 13; DB 1; Length 12;  
Best Local Similarity 33.3%; Pred. No. 4e+03;  
Matches 1; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

QY 4 WXF 6  
|  
Db 8 WKY 10

RESULT 11  
UR2B\_CATCO  
ID UR2B CATCO STANDARD; PRT; 12 AA.  
AC P04559;  
DT 13-AUG-1987 (Rel. 05, Created)  
DT 13-AUG-1987 (Rel. 05, Last sequence update)  
DT 16-OCT-2001 (Rel. 40, Last annotation update)  
DE Urotensin IIB (U-IIB) (UIIB).  
OS Catostomus commersoni (White sucker).  
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
OC Actinopterygii; Neopterygii; Teleostei; Ostariophysi; Cypriniformes;  
OC Catostomidae; Catostomus.  
OX NCBI\_TaxID=7971;  
RN [1]  
RP SEQUENCE.  
RX MEDLINE=84041959; PubMed=6138758;  
RA McMaster D., Lederis K.;  
RT "Isolation and amino acid sequence of two urotensin II peptides from  
RT Catostomus commersoni urophyses.";  
RL Peptides 4:367-373(1983).  
CC -!- FUNCTION: UROTENSIN IS FOUND IN THE TELEOST CAUDAL NEUROSECRETORY  
CC SYSTEM. IT HAS A SUGGESTED ROLE IN OSMOREGULATION AND AS A  
CC CORTICOTROPIN-RELEASING FACTOR.

Query Match 41.9%; Score 13; DB 1; Length 12;  
Best Local Similarity 33.3%; Pred. No. 4e+03;  
Matches 1; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

QY 4 WXF 6  
|  
Db 8 WKY 10

RESULT 12  
UR2B\_CYPCA  
ID UR2B CYPCA STANDARD; PRT; 12 AA.  
AC P04561;  
DT 13-AUG-1987 (Rel. 05, Created)  
DT 13-AUG-1987 (Rel. 05, Last sequence update)  
DT 16-OCT-2001 (Rel. 40, Last annotation update)  
DE Urotensin II-beta.  
OS Cyprinus carpio (Common carp).  
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
OC Actinopterygii; Neopterygii; Teleostei; Ostariophysi; Cypriniformes;  
OC Cyprinidae; Cyprinus.  
OX NCBI\_TaxID=7962;  
RN [1]

RP SEQUENCE.  
RA Muneakata E., Ohtaki T., Ichikawa T., McMaster D., Lederis K.;  
RL (in) Rich D.H., Gross E. (eds.);  
RL Proceedings of the 7th american peptide symposium, pp.69-72,  
RL Pierce Chemical Co., Rockford IL. (1981).  
CC -!- FUNCTION: UROTENSIN IS FOUND IN THE TELEOST CAUDAL NEUROSECRETORY  
CC SYSTEM. IT HAS A SUGGESTED ROLE IN OSMOREGULATION AND AS A  
CC CORTICOTROPIN-RELEASING FACTOR.  
CC -!- SIMILARITY: BELONGS TO THE UROTENSIN 2 FAMILY.  
DR InterPro; IPR001483; Urotensin\_II.  
DR Pfam; PF02083; Urotensin\_II; 1.  
DR PROSITE; PS00984; UROTENSIN\_II; 1.  
KW Hormone.  
FT DISULFID 6 11  
FT VARIANT 2 2 G -> S.  
SQ SEQUENCE 12 AA; 1407 MW; 73960A9FB879CEBB CRC64;  
  
Query Match 41.9%; Score 13; DB 1; Length 12;  
Best Local Similarity 33.3%; Pred. No. 4e+03;  
Matches 1; Conservative 1; Mismatches 1; Indels 0; Gaps 0;  
  
QY 4 WXF 6  
Db 8 WKY 10  
  
RESULT 13  
UR2\_GILMI  
ID UR2\_GILMI STANDARD; PRT; 12 AA.  
AC P01147;  
DT 21-JUL-1986 (Rel. 01, Created)  
DT 21-JUL-1986 (Rel. 01, Last sequence update)  
DT 16-OCT-2001 (Rel. 40, Last annotation update)  
DE Urotensin II (U-II) (UII).  
OS Gillichthys mirabilis (Long-jawed mudsucker).  
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
OC Actinopterygii; Neopterygii; Teleostei; Euteleostei; Neoteleostei;  
OC Acanthomorpha; Acanthopterygii; Percomorpha; Perciformes; Gobioidae;  
OC Gobiidae; Gillichthys.  
OX NCBI\_TaxID=8222;  
RN [1]  
RP SEQUENCE.  
RX MEDLINE=81054904; PubMed=6107911;  
RA Pearson D., Shively J.E., Clark B.R., Geschwind I.I., Barkley M.,  
RA Nishioka R., Bern H.A.;  
RT "Urotensin II: a somatostatin-like peptide in the caudal  
RT neurosecretory system of fishes."  
RL Proc. Natl. Acad. Sci. U.S.A. 77:5021-5024(1980).  
CC -!- FUNCTION: UROTENSIN IS FOUND IN THE TELEOST CAUDAL NEUROSECRETORY  
CC SYSTEM. IT HAS A SUGGESTED ROLE IN OSMOREGULATION AND AS A  
CC CORTICOTROPIN-RELEASING FACTOR.  
CC -!- SIMILARITY: BELONGS TO THE UROTENSIN 2 FAMILY.  
DR PIR; A01409; UOGM2.  
DR PIR; S42765; S42765.  
DR InterPro; IPR001483; Urotensin\_II.  
DR Pfam; PF02083; Urotensin\_II; 1.  
DR PROSITE; PS00984; UROTENSIN\_II; 1.  
KW Hormone.  
FT DISULFID 6 11  
SQ SEQUENCE 12 AA; 1364 MW; 968BF8982679CEBA CRC64;  
  
Query Match 41.9%; Score 13; DB 1; Length 12;  
Best Local Similarity 33.3%; Pred. No. 4e+03;  
Matches 1; Conservative 1; Mismatches 1; Indels 0; Gaps 0;  
  
QY 4 WXF 6  
Db 8 WKY 10  
  
RESULT 14  
UR2\_POLSP  
ID UR2\_POLSP STANDARD; PRT; 12 AA.

AC P81022;  
DT 01-NOV-1997 (Rel. 35, Created)  
DT 01-NOV-1997 (Rel. 35, Last sequence update)  
DT 16-OCT-2001 (Rel. 40, Last annotation update)  
DE Urotensin II (U-II) (UII).  
OS Polyodon spathula (North American paddlefish).  
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
OC Actinopterygii; Chondrostei; Acipenseriformes; Polyodontidae;  
OC Polyodon.  
OX NCBI\_TaxID=7913;  
RN [1]  
RP SEQUENCE FROM N.A.  
RX TISSUE=Spinal cord;  
RX MEDLINE=96051494; PubMed=8536944;  
RA Waugh D., Youson J., Mims S.D., Sower S., Conlon J.M.;  
RT "Urotensin II from the river lamprey (Lampetra fluviatilis), the sea  
RT lamprey (Petromyzon marinus), and the paddlefish (Polyodon  
RT spathula).";  
RL Gen. Comp. Endocrinol. 99:323-332(1995).  
CC -!- FUNCTION: HAS A SUGGESTED ROLE IN OSMOREGULATION AND AS A  
CC CORTICOTROPIN-RELEASING FACTOR. PROBABLY INVOLVED IN SMOOTH  
CC MUSCLE STIMULATION.  
CC -!- SIMILARITY: BELONGS TO THE UROTENSIN 2 FAMILY.  
DR InterPro; IPR001483; Urotensin\_II.  
DR Pfam; PF02083; Urotensin\_II; 1.  
DR PROSITE; PS00984; UROTENSIN\_II; 1.  
KW Hormone.  
FT DISULFID 6 11 BY SIMILARITY.  
SQ SEQUENCE 12 AA; 1410 MW; 7551E9DBB879CEBB CRC64;  
  
Query Match 41.9%; Score 13; DB 1; Length 12;  
Best Local Similarity 33.3%; Pred. No. 4e+03;  
Matches 1; Conservative 1; Mismatches 1; Indels 0; Gaps 0;  
  
QY 4 WXF 6  
Db 8 WKY 10  
  
RESULT 15  
UR2\_SCYCA  
ID UR2\_SCYCA STANDARD; PRT; 12 AA.  
AC P35490;  
DT 01-JUN-1994 (Rel. 29, Created)  
DT 01-JUN-1994 (Rel. 29, Last sequence update)  
DT 16-OCT-2001 (Rel. 40, Last annotation update)  
DE Urotensin II (U-II) (UII).  
OS Scyliorhinus canicula (Spotted dogfish) (Spotted catshark).  
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Chondrichthyes;  
OC Elasmobranchii; Galeomorphii; Galeoidea; Carcharhiniformes;  
OC Scyliorhinidae; Scyliorhinus.  
OX NCBI\_TaxID=7830;  
RN [1]  
RP SEQUENCE.  
RX TISSUE=Spinal cord;  
RX MEDLINE=92319231; PubMed=1620290;  
RA Conlon J.M., O'Harte F., Smith D.D., Balment R.J., Hazon N.;  
RT "Purification and characterization of urotensin II and parvalbumin  
RT from an elasmobranch fish, Scyliorhinus canicula (common dogfish).";  
RL Neuroendocrinology 55:230-235(1992).  
CC -!- FUNCTION: HAS A SUGGESTED ROLE IN OSMOREGULATION AND AS A  
CC CORTICOTROPIN-RELEASING FACTOR. PROBABLY INVOLVED IN SMOOTH  
CC MUSCLE STIMULATION.  
CC -!- SIMILARITY: BELONGS TO THE UROTENSIN 2 FAMILY.  
DR InterPro; IPR001483; Urotensin\_II.  
DR Pfam; PF02083; Urotensin\_II; 1.  
DR PROSITE; PS00984; UROTENSIN\_II; 1.  
KW Hormone.  
FT DISULFID 6 11  
SQ SEQUENCE 12 AA; 1526 MW; 804729F9D579CEBA CRC64;  
  
Query Match 41.9%; Score 13; DB 1; Length 12;  
Best Local Similarity 33.3%; Pred. No. 4e+03;

Matches 1; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

Oy 4 WXF 6

| :

Db 8 WKY 10

Search completed: December 3, 2003, 11:51:51  
Job time : 7.33333 secs



GenCore version 5.1.6  
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OM protein - protein search, using sw model

Run on: December 3, 2003, 11:48:35 ; Search time 11 Seconds  
(without alignments)  
52.456 Million cell updates/sec

Title: US-09-912-414-9  
Perfect score: 31  
Sequence: 1 WXXWXF 6  
Scoring table: BLOSUM62  
Gapop 10.0 , Gapext 0.5

Searched: 283308 seqs, 96168682 residues

Total number of hits satisfying chosen parameters: 2520

Minimum DB seq length: 0  
Maximum DB seq length: 15

Post-processing: Minimum Match 0%  
Maximum Match 100%  
Listing first 45 summaries

Database : PIR\_76:\*  
1: pir1:\*  
2: pir2:\*  
3: pir3:\*  
4: pir4:\*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Query			DB ID	Description
	Score	Match	Length		
1	21	67.7	9	2 A43848	cell surface adhes
2	20	64.5	10	2 F49033	T-cell receptor ga
3	20	64.5	12	2 PH1324	Ig heavy chain DJ
4	20	64.5	12	2 PH1308	Ig heavy chain DJ
5	20	64.5	13	2 S61798	T-cell-specific tr
6	20	64.5	14	2 PH1322	Ig heavy chain DJ
7	17	54.8	13	2 S23372	T-cell receptor al
8	17	54.8	13	2 B25448	Ig kappa-1 chain,
9	17	54.8	13	2 B26406	Ig kappa chain J r
10	17	54.8	13	2 A47630	Ig kappa chain J r
11	16	51.6	8	2 T13818	cytochrome oxidase
12	16	51.6	10	2 T17054	cytochrome-c oxida
13	16	51.6	10	2 T13976	cytochrome-c oxida
14	16	51.6	10	2 T17057	cytochrome-c oxida
15	16	51.6	10	2 T12303	cytochrome-c oxida
16	16	51.6	10	2 T14019	cytochrome-c oxida
17	16	51.6	10	2 T17060	cytochrome-c oxida
18	16	51.6	10	2 T17063	cytochrome-c oxida
19	16	51.6	10	2 T12325	cytochrome-c oxida
20	16	51.6	10	2 T14043	cytochrome-c oxida
21	16	51.6	10	2 T14054	cytochrome-c oxida
22	16	51.6	10	2 T17066	cytochrome-c oxida
23	16	51.6	10	2 T17069	cytochrome-c oxida
24	16	51.6	10	2 T12308	cytochrome-c oxida
25	16	51.6	10	2 T17072	cytochrome-c oxida
26	16	51.6	10	2 T12312	cytochrome-c oxida
27	16	51.6	10	2 T12329	cytochrome-c oxida
28	16	51.6	10	2 T12316	cytochrome-c oxida
29	16	51.6	10	2 T12321	cytochrome-c oxida

30	16	51.6	10	2 T14215	cytochrome-c oxida
31	16	51.6	10	2 T14223	cytochrome-c oxida
32	16	51.6	10	2 T14219	cytochrome-c oxida
33	16	51.6	12	2 A29169	phospholipase A2 (
34	16	51.6	14	2 PT0077	proteochonditin c
35	16	51.6	15	2 PA0099	phenotypic variati
36	15	48.4	9	2 S56004	glucan 1,3-beta-gl
37	15	48.4	15	2 S24159	leukocyte elastase
38	14	45.2	9	2 S07241	litorin - Rohde's
39	14	45.2	10	1 RHFGG	gonadoliberin - pi
40	14	45.2	10	1 RHSHG	gonadoliberin - sh
41	14	45.2	10	1 RHAQ1	gonadoliberin I -
42	14	45.2	10	2 A21114	gonadoliberin - ch
43	14	45.2	11	2 S68649	spermadhesin AQN-3
44	14	45.2	15	2 PH1365	Ig heavy chain DJ
45	13	41.9	9	2 S07205	litorin 2-Glu - Au

ALIGNMENTS

RESULT 1

A43848  
cell surface adhesin for heparan sulfate, 66K - Staphylococcus aureus (fragment)  
C;Species: Staphylococcus aureus  
C;Date: 10-Mar-1993 #sequence\_revision 18-Nov-1994 #text\_change 24-Feb-1995  
C;Accession: A43848  
R;Liang, O.D.; Ascencio, F.; Fransson, L.A.; Wadstrom, T.  
Infect. Immun. 60, 899-906, 1992  
A;Title: Binding of heparan sulfate to Staphylococcus aureus.  
A;Reference number: A43848; MUID:92176005; PMID:1541563  
A;Accession: A43848  
A;Status: preliminary  
A;Molecule type: protein  
A;Residues: 1-9 <LIA>  
A;Note: sequence extracted from NCBI backbone (NCBIP:85442)

Query Match 67.7%; Score 21; DB 2; Length 9;  
Best Local Similarity 50.0%; Pred. No. 2.8e+05;  
Matches 2; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

Qy 1 WXXW 4  
Db 2 WTGW 5

RESULT 2

F49033  
T-cell receptor gamma chain V-D-J region - human (fragment)  
C;Species: Homo sapiens (man)  
C;Date: 19-Dec-1993 #sequence\_revision 17-Mar-2000 #text\_change 17-Mar-2000  
C;Accession: F49033  
R;Morita, C.T.; Verma, S.; Aparicio, P.; Martinez, C.; Spits, H.; Brenner, M.B.  
Eur. J. Immunol. 21, 2999-3007, 1991  
A;Title: Functionally distinct subsets of human gamma/delta T cells.  
A;Reference number: A49033; MUID:92083926; PMID:1684157  
A;Accession: F49033  
A;Status: preliminary  
A;Molecule type: DNA  
A;Residues: 1-10 <MOR>  
A;Cross-references: GB:S72605; NID:G240700; PIDN:AAB20632.1; PID:G240701  
A;Note: sequence extracted from NCBI backbone (NCBIN:72605, NCBIP:72606)  
C;Keywords: T-cell receptor

Query Match 64.5%; Score 20; DB 2; Length 10;  
Best Local Similarity 50.0%; Pred. No. 5.7e+02;  
Matches 2; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

Qy 1 WXXW 4  
Db 4 WERW 7

## RESULT 3

PH1324  
Ig heavy chain DJ region (clone C510-100) - human (fragment)  
C;Species: Homo sapiens (man)  
C;Date: 30-Sep-1993 #sequence\_revision 30-Sep-1993 #text\_change 07-May-1999  
C;Accession: PH1324  
R;Wasserman, R.; Galili, N.; Ito, Y.; Reichard, B.A.; Shane, S.; Rovera, G.  
J. Exp. Med. 176, 1577-1581, 1992  
A;Title: Predominance of fetal type DJH joining in young children with B precursor lymphoma  
A;Reference number: PH1302; MUID:93094761; PMID:1460419  
A;Accession: PH1324  
A;Molecule type: DNA  
A;Residues: 1-12 <WAS>  
C;Keywords: heterotetramer; immunoglobulin

Query Match 64.5%; Score 20; DB 2; Length 12;  
Best Local Similarity 50.0%; Pred. No. 6.6e+02;  
Matches 2; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

Qy 1 WXXW 4  
| |  
Db 5 WYVW 8

## RESULT 4

PH1308  
Ig heavy chain DJ region (clone C731-94) - human (fragment)  
C;Species: Homo sapiens (man)  
C;Date: 30-Sep-1993 #sequence\_revision 30-Sep-1993 #text\_change 07-May-1999  
C;Accession: PH1308  
R;Wasserman, R.; Galili, N.; Ito, Y.; Reichard, B.A.; Shane, S.; Rovera, G.  
J. Exp. Med. 176, 1577-1581, 1992  
A;Title: Predominance of fetal type DJH joining in young children with B precursor lymphoma  
A;Reference number: PH1302; MUID:93094761; PMID:1460419  
A;Accession: PH1308  
A;Molecule type: DNA  
A;Residues: 1-12 <WAS>  
C;Keywords: heterotetramer; immunoglobulin

Query Match 64.5%; Score 20; DB 2; Length 12;  
Best Local Similarity 50.0%; Pred. No. 6.6e+02;  
Matches 2; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

Qy 1 WXXW 4  
| |  
Db 7 WQVW 10

## RESULT 5

S61798  
T-cell-specific transcription factor 1 splice form G - human (fragment)  
N;Alternate names: transcription factor TCF-1g  
C;Species: Homo sapiens (man)  
C;Date: 19-Mar-1997 #sequence\_revision 18-Jul-1997 #text\_change 24-Jul-1998  
C;Accession: S61798; S61880  
R;Mayer, K.; Wolff, E.; Clevers, H.; Ballhausen, W.G.  
Biochim. Biophys. Acta 1263, 169-172, 1995  
A;Title: The human high mobility group (HMG)-box transcription factor TCF-1: novel isoform  
A;Reference number: S61796; MUID:95367594; PMID:7640309  
A;Accession: S61798  
A;Molecule type: mRNA  
A;Residues: 1-13 <MAY>  
A;Cross-references: EMBL:Z47364  
A;Note: DNA was also sequenced  
C;Keywords: alternative splicing; DNA binding; transcription factor

Query Match 64.5%; Score 20; DB 2; Length 13;  
Best Local Similarity 50.0%; Pred. No. 7e+02;  
Matches 2; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

Qy 1 WXXW 4  
| |  
Db 6 WQVW 9

## RESULT 6

PH1322  
Ig heavy chain DJ region (clone C344-99) - human (fragment)  
C;Species: Homo sapiens (man)  
C;Date: 30-Sep-1993 #sequence\_revision 30-Sep-1993 #text\_change 07-May-1999  
C;Accession: PH1322  
R;Wasserman, R.; Galili, N.; Ito, Y.; Reichard, B.A.; Shane, S.; Rovera, G.  
J. Exp. Med. 176, 1577-1581, 1992  
A;Title: Predominance of fetal type DJH joining in young children with B precursor lymphoma  
A;Reference number: PH1302; MUID:93094761; PMID:1460419  
A;Accession: PH1322  
A;Molecule type: DNA  
A;Residues: 1-14 <WAS>  
C;Keywords: heterotetramer; immunoglobulin

Query Match 64.5%; Score 20; DB 2; Length 14;  
Best Local Similarity 50.0%; Pred. No. 7.4e+02;  
Matches 2; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

Qy 1 WXXW 4  
| |  
Db 6 WYVW 9

## RESULT 7

S23372  
T-cell receptor alpha chain J region - human (fragment)  
C;Species: Homo sapiens (man)  
C;Date: 22-Nov-1993 #sequence\_revision 26-May-1995 #text\_change 17-Mar-1999  
C;Accession: S23372  
R;Pluschke, G.; Ricken, G.; Taube, H.; Kroninger, S.; Melchers, I.; Peter, H.H.; Eichmeier, J. Immunol. 21, 2749-2754, 1991  
A;Title: Biased T cell receptor V(alpha) region repertoire in the synovial fluid of rheumatoid arthritis  
A;Reference number: S23364; MUID:92037820; PMID:1657615  
A;Accession: S23372  
A;Status: preliminary; translation not shown  
A;Molecule type: mRNA  
A;Residues: 1-13 <PLU>  
A;Cross-references: EMBL:X58167  
C;Keywords: T-cell receptor

Query Match 54.8%; Score 17; DB 2; Length 13;  
Best Local Similarity 66.7%; Pred. No. 2.2e+03;  
Matches 2; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

Qy 4 WXF 6  
| |  
Db 11 WTF 13

## RESULT 8

B25448  
Ig kappa-1 chain, 69 allotype, J-KI.1 segment - rabbit (fragment)  
C;Species: Oryctolagus cuniculus (domestic rabbit)  
C;Date: 16-Aug-1988 #sequence\_revision 16-Aug-1988 #text\_change 05-Nov-1999  
C;Accession: B25448  
R;Akimenko, M.A.; Mariame, B.; Rougeon, F.  
Proc. Natl. Acad. Sci. U.S.A. 83, 5180-5183, 1986  
A;Title: Evolution of the immunoglobulin kappa light chain locus in the rabbit: evidence for a gene conversion event  
A;Reference number: A94110; MUID:86259753; PMID:3088570  
A;Accession: B25448  
A;Molecule type: DNA  
A;Residues: 1-13 <AKI>  
A;Cross-references: GB:M14067; GB:M14062; GB:M14063; GB:M14064; GB:M14065; GB:M14066; I  
C;Keywords: heterotetramer; immunoglobulin

Query Match 54.8%; Score 17; DB 2; Length 13;  
Best Local Similarity 66.7%; Pred. No. 2.2e+03;  
Matches 2; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

Qy 4 WXF 6

Db                   |||  
                    1 WAF 3

RESULT 9  
B26406  
Ig kappa chain J region - mouse  
C;Species: Mus musculus (house mouse)  
C;Date: 30-Jun-1989 #sequence\_revision 30-Jun-1989 #text\_change 16-Aug-1996  
C;Accession: B26406  
R;Sanz, I.; Capra, J.D.  
Proc. Natl. Acad. Sci. U.S.A. 84, 1085-1089, 1987  
A;Title: V-K and J-K gene segments of A/J Ars-A antibodies: somatic recombination genera  
A;Reference number: A26406; MUID:87147197; PMID:3103124  
A;Accession: B26406  
A;Molecule type: DNA  
A;Residues: 1-13 <SAN>  
A;Cross-references: GB:M15519  
C;Keywords: heterotetramer; immunoglobulin

Query Match           54.8%; Score 17; DB 2; Length 13;  
Best Local Similarity   66.7%; Pred. No. 2.2e+03;  
Matches   2; Conservative   0; Mismatches   1; Indels   0; Gaps   0;

QY                   4 WXF 6  
                    |||  
Db                   1 WTF 3

RESULT 10  
A47630  
Ig kappa chain-J region J1 - southeastern Australian rat  
C;Species: Rattus sordidus villosissimus (southeastern Australian rat)  
C;Date: 03-Feb-1994 #sequence\_revision 03-Feb-1994 #text\_change 05-Nov-1999  
C;Accession: A47630  
R;Gutman, G.A.; Besta, R.M.; Frank, M.B.; Baverstock, P.R.  
Immunogenetics 26, 14-20, 1987  
A;Title: Duplication of J-kappa genes within genus Rattus.  
A;Reference number: A47630; MUID:87278355; PMID:3111993  
A;Accession: A47630  
A;Status: preliminary; not compared with conceptual translation  
A;Molecule type: DNA  
A;Residues: 1-13 <GUT>  
A;Cross-references: GB:M17319; NID:g204788; PIDN:AAA41397.1; PID:g204789  
C;Keywords: heterotetramer; immunoglobulin

Query Match           54.8%; Score 17; DB 2; Length 13;  
Best Local Similarity   66.7%; Pred. No. 2.2e+03;  
Matches   2; Conservative   0; Mismatches   1; Indels   0; Gaps   0;

QY                   4 WXF 6  
                    |||  
Db                   1 WTF 3

RESULT 11  
T13818  
cytochrome oxidase subunit I - Atlantic hagfish mitochondrion (fragment)  
C;Species: mitochondrion Myxine glutinosa (Atlantic hagfish)  
C;Date: 20-Sep-1999 #sequence\_revision 20-Sep-1999 #text\_change 21-Jul-2000  
C;Accession: T13818  
R;Delarbre, C.; Barriol, V.; Tillier, S.; Janvier, P.; Gachelin, G.  
Mol. Biol. Evol. 14, 807-813, 1997  
A;Title: The main features of the craniate mitochondrial DNA between the ND1 and the COI  
A;Reference number: Z17775; MUID:97398704; PMID:9254918  
A;Accession: T13818  
A;Status: preliminary; translated from GB/EMBL/DBJ  
A;Molecule type: DNA  
A;Residues: 1-8 <DEL>  
A;Cross-references: EMBL:Y09527; NID:g2340019; PIDN:CAA70718.1; PID:g2340022  
C;Genetics:  
A;Genome: mitochondrion  
A;Note: COI

C;Keywords: mitochondrion

Query Match           51.6%; Score 16; DB 2; Length 8;  
Best Local Similarity   66.7%; Pred. No. 2.8e+05;  
Matches   2; Conservative   0; Mismatches   1; Indels   0; Gaps   0;

QY                   4 WXF 6  
                    |||  
Db                   6 WFF 8

RESULT 12  
T17054  
cytochrome-c oxidase (EC 1.9.3.1) chain I - Basiliscus plumifrons mitochondrion (fragme  
C;Species: mitochondrion Basiliscus plumifrons  
C;Date: 15-Oct-1999 #sequence\_revision 15-Oct-1999 #text\_change 22-Oct-1999  
C;Accession: T17054  
R;Macey, J.R.; Larson, A.; Ananjeva, N.B.; Papenfuss, T.J.  
J. Mol. Evol. 44, 660-674, 1997  
A;Title: Evolutionary shifts in three major structural features of the mitochondrial g  
A;Reference number: Z18674; MUID:97315309; PMID:9169559  
A;Accession: T17054  
A;Status: preliminary; translated from GB/EMBL/DBJ  
A;Molecule type: DNA  
A;Residues: 1-10 <MAC>  
A;Cross-references: EMBL:U82680; NID:g3603104; PID:g3603107; PIDN:AAC62269.1  
C;Genetics:  
A;Genome: mitochondrion  
A;Note: COI  
C;Keywords: mitochondrion; oxidoreductase

Query Match           51.6%; Score 16; DB 2; Length 10;  
Best Local Similarity   66.7%; Pred. No. 2.6e+03;  
Matches   2; Conservative   0; Mismatches   1; Indels   0; Gaps   0;

QY                   4 WXF 6  
                    |||  
Db                   6 WLF 8

RESULT 13  
T13976  
cytochrome-c oxidase (EC 1.9.3.1) chain I - Cnemidophorus tigris mitochondrion (fragme;  
C;Species: mitochondrion Cnemidophorus tigris  
C;Date: 20-Sep-1999 #sequence\_revision 20-Sep-1999 #text\_change 11-May-2000  
C;Accession: T13976  
R;Macey, J.R.; Larson, A.; Ananjeva, N.B.; Fang, Z.; Papenfuss, T.J.  
Mol. Biol. Evol. 14, 91-104, 1997  
A;Title: Two novel gene orders and the role of light-strand replication in rearrangeme;  
A;Reference number: Z17789; MUID:97153826; PMID:9000757  
A;Accession: T13976  
A;Status: preliminary; translated from GB/EMBL/DBJ  
A;Molecule type: DNA  
A;Residues: 1-10 <MAC>  
A;Cross-references: EMBL:U71332; NID:g1753236; PID:g1753239; PIDN:AAB48274.1  
C;Genetics:  
A;Genome: mitochondrion  
A;Note: COI  
C;Keywords: mitochondrion; oxidoreductase

Query Match           51.6%; Score 16; DB 2; Length 10;  
Best Local Similarity   66.7%; Pred. No. 2.6e+03;  
Matches   2; Conservative   0; Mismatches   1; Indels   0; Gaps   0;

QY                   4 WXF 6  
                    |||  
Db                   6 WFF 8

RESULT 14  
T17057  
cytochrome-c oxidase (EC 1.9.3.1) chain I - Crotaphytus collaris mitochondrion (fragme;  
C;Species: mitochondrion Crotaphytus collaris

C;Date: 15-Oct-1999 #sequence\_revision 15-Oct-1999 #text\_change 22-Oct-1999  
C;Accession: T17057  
R;Macey, J.R.; Larson, A.; Ananjeva, N.B.; Papenfuss, T.J.  
J. Mol. Evol. 44, 660-674, 1997  
A;Title: Evolutionary shifts in three major structural features of the mitochondrial gen  
A;Reference number: Z18674; MUID:97315309; PMID:9169559  
A;Accession: T17057  
A;Status: preliminary; translated from GB/EMBL/DBJ  
A;Molecule type: DNA  
A;Residues: 1-10 <MAC>  
A;Cross-references: EMBL:U82681; NID:g3603108; PID:g3603111; PIDN:AAC62272.1  
C;Genetics:  
A;Genome: mitochondrion  
A;Note: COI  
C;Keywords: mitochondrion; oxidoreductase

Query Match 51.6%; Score 16; DB 2; Length 10;  
Best Local Similarity 66.7%; Pred. No. 2.6e+03;  
Matches 2; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

QY 4 WXF 6  
| |  
Db 6 WFF 8

RESULT 15  
T12303  
cytochrome-c oxidase (EC 1.9.3.1) chain I - Diposaurus dorsalis mitochondrion (fragment  
C;Species: mitochondrion Diposaurus dorsalis  
C;Date: 23-Jul-1999 #sequence\_revision 23-Jul-1999 #text\_change 22-Oct-1999  
C;Accession: T12303  
R;Schulte, J.A.; Macey, J.R.; Larson, A.; Papenfuss, T.J.  
Mol. Phylogenet. Evol. 10, 367-376, 1998  
A;Title: Molecular tests of phylogenetic taxonomies: A general procedure and example usi  
A;Reference number: Z17488; MUID:99162288; PMID:10051389  
A;Accession: T12303  
A;Status: preliminary; translated from GB/EMBL/DBJ  
A;Molecule type: DNA  
A;Residues: 1-10 <SCH>  
A;Cross-references: EMBL:AF049857; NID:g4105726; PID:g4105729; PIDN:AAD02514.1  
C;Genetics:  
A;Genome: mitochondrion  
A;Note: COI  
C;Keywords: mitochondrion; oxidoreductase

Query Match 51.6%; Score 16; DB 2; Length 10;  
Best Local Similarity 66.7%; Pred. No. 2.6e+03;  
Matches 2; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

QY 4 WXF 6  
| |  
Db 6 WFF 8

Search completed: December 3, 2003, 11:54:08  
Job time : 11 secs

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OM protein - protein search, using sw model

Run on: December 12, 2003, 10:26:30 ; Search time 30.3 Seconds  
(without alignments)  
31.431 Million cell updates/sec

Title: US-09-912-414-9  
Perfect score: 31  
Sequence: 1 WXXWXF 6

Scoring table: BLOSUM62  
Gapop 10.0 , Gapext 0.5

Searched: 1107863 seqs, 158726573 residues

Total number of hits satisfying chosen parameters: 350435

Minimum DB seq length: 0  
Maximum DB seq length: 15

Post-processing: Minimum Match 0%  
Maximum Match 100%  
Listing first 45 summaries

Database : A Geneseq 19Jun03:\*  
1: /SIDS1/gcgdata/geneseq/geneseq-emb1/AA1980.DAT:\*  
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3: /SIDS1/gcgdata/geneseq/geneseq-emb1/AA1982.DAT:\*  
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23: /SIDS1/gcgdata/geneseq/geneseq-emb1/AA2002.DAT:\*  
24: /SIDS1/gcgdata/geneseq/geneseq-emb1/AA2003.DAT:\*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	ID	Description
1	28	90.3	15	AA1980	Epitope derived fr
2	28	90.3	15	AA1981	Streptococcus pneu
3	27	87.1	9	AA26751	Fibrin binding loo
4	27	87.1	15	AA26733	Fibrin binding pep
5	26	83.9	6	AA1505	Peptide which bind
6	26	83.9	6	AA1506	Peptide which bind
7	26	83.9	6	AA1508	Peptide which bind
8	26	83.9	6	AA1513	Staphylococcus aur
9	26	83.9	6	AA1514	Staphylococcus aur

10	26	83.9	6	24	ABR45369	Staphylococcus aur
11	26	83.9	6	24	ABR45370	Staphylococcus aur
12	26	83.9	6	24	ABR45425	Staphylococcus aur
13	26	83.9	6	24	ABR45426	Staphylococcus aur
14	26	83.9	6	24	ABR45481	Staphylococcus aur
15	26	83.9	6	24	ABR45482	Staphylococcus aur
16	26	83.9	6	24	ABR45593	Staphylococcus aur
17	26	83.9	6	24	ABR45594	Staphylococcus aur
18	26	83.9	9	23	AAE26775	Fibrin binding pep
19	26	83.9	15	21	AAE65508	Oestrogen receptor
20	26	83.9	15	23	ABB99042	Serine/threonine p
21	26	83.9	15	23	AAE26759	Fibrin binding pep
22	26	83.9	15	23	AAU86245	Oestrogen receptor
23	25	80.6	6	15	AAE57391	Peptide for treati
24	25	80.6	6	21	AAE01492	Peptide which bind
25	25	80.6	6	21	AAE01497	Peptide which bind
26	25	80.6	6	21	AAE01499	Peptide which bind
27	25	80.6	6	24	ABR44865	Staphylococcus aur
28	25	80.6	6	24	ABR44866	Staphylococcus aur
29	25	80.6	6	24	ABR45311	Staphylococcus aur
30	25	80.6	6	24	ABR45312	Staphylococcus aur
31	25	80.6	6	24	ABR45367	Staphylococcus aur
32	25	80.6	6	24	ABR45368	Staphylococcus aur
33	25	80.6	6	24	ABR45423	Staphylococcus aur
34	25	80.6	6	24	ABR45424	Staphylococcus aur
35	25	80.6	6	24	ABR45479	Staphylococcus aur
36	25	80.6	6	24	ABR45480	Staphylococcus aur
37	25	80.6	6	24	ABR45537	Staphylococcus aur
38	25	80.6	6	24	ABR45538	Staphylococcus aur
39	25	80.6	6	24	ABR45591	Staphylococcus aur
40	25	80.6	6	24	ABR45592	Staphylococcus aur
41	25	80.6	7	22	AAE45777	H11 binding site c
42	25	80.6	9	21	AAE01498	Peptide which bind
43	25	80.6	11	21	AAE20714	Polymeric immunogl
44	25	80.6	13	18	AAE38112	Dystrophin WW doma
45	25	80.6	14	22	AAE07760	Human HLA-DP restr

ALIGNMENTS

RESULT 1

AA1980  
ID AAY30351 standard; Peptide; 15 AA.

XX  
AC AAY30351;

XX  
DT 09-NOV-1999 (first entry)

XX  
DE Epitope derived from pneumococcal surface adhesion A protein.

XX  
KW Pneumococcal surface adhesion A protein; PsaA; monoclonal antibody;  
vaccine; Streptococcus pneumoniae infection.

XX  
OS Streptococcus pneumoniae.

XX  
PN WO9945121-A1.

XX  
PD 10-SEP-1999.

XX  
PF 26-FEB-1999; 99WO-US04326.

XX  
PR 02-MAR-1998; 98US-0076565.

XX  
PA (USSH ) US DEPT HEALTH & HUMAN SERVICES.

XX  
PI Ades EW, Carlone GM, Sampson JS, Tharpe JA, Westerink MAJ;

XX  
PI Zeiler JL;

XX  
XX WPI; 1999-540849/45.

XX  
DR New peptides corresponding to Streptococcus pneumoniae PsaA, used  
for treating or preventing Streptococcus pneumoniae infection in a

XX  
PT



PT subject  
XX Claim 6; Page 43; 58pp; English.  
PS  
XX  
CC AAY30351-54 represent immunogenic peptides which are derived from  
CC a pneumococcal surface adhesion A protein (PsaA). The specification  
CC describes monoclonal antibodies which bind epitopes of the PsaA protein  
CC (e.g present sequence). The peptides can be used in vaccines to prevent  
CC Streptococcus pneumoniae infections. The antibodies of the invention  
CC can also be used to detect S. pneumoniae in a sample or individual.  
XX  
SQ Sequence 15 AA;  
  
Query Match 90.3%; Score 28; DB 20; Length 15;  
Best Local Similarity 50.0%; Pred. No. 1.1e+02;  
Matches 3; Conservative 0; Mismatches 3; Indels 0; Gaps 0;  
  
QY 1 WXXWXF 6  
| | |  
Db 7 WTAWAF 12  
  
RESULT 2  
AAE19239  
ID AAE19239 standard; peptide; 15 AA.  
XX  
AC AAE19239;  
XX  
DT 21-MAY-2002 (first entry)  
XX  
DE Streptococcus pneumoniae PsaA immunogenic peptide #1.  
XX  
KW Multiple antigenic peptide; MAP; immunogenic; immunity; infection;  
KW pneumococcal surface adhesin protein A; PsaA; antibacterial.  
XX  
OS Streptococcus pneumoniae.  
XX WO200204497-A2.  
PN  
XX  
PD 17-JAN-2002.  
XX  
PF 10-JUL-2001; 2001WO-US21626.  
XX  
PR 10-JUL-2000; 2000US-0613092.  
XX  
PA (USSH ) US DEPT HEALTH & HUMAN SERVICES.  
XX  
PI Ades EW, Johnson SE, Jue DL, Sampson JS, Carlone GM;  
XX  
DR WPI; 2002-195762/25.  
XX  
PT New multiple antigenic peptide for immunizing against streptococcal  
PT infections, binds to monoclonal antibody obtained in response to  
PT immunizing an animal with pneumococcal surface adhesion protein A or  
PT its fragment  
XX  
PS Claim 2; Page 56; 86pp; English.  
XX  
CC The invention relates to multiple antigenic peptides (MAP) immunogenic  
CC against Streptococcus pneumoniae. MAP binds to monoclonal antibody  
CC obtained in response to immunising an animal with pneumococcal surface  
CC adhesion protein A (PsaA) or its fragment. MAP is useful for conferring  
CC protective immunity against S. pneumoniae infection in a subject. The  
CC present sequence is Streptococcus pneumoniae PsaA immunogenic peptide.  
XX  
SQ Sequence 15 AA;  
  
Query Match 90.3%; Score 28; DB 23; Length 15;  
Best Local Similarity 50.0%; Pred. No. 1.1e+02;  
Matches 3; Conservative 0; Mismatches 3; Indels 0; Gaps 0;  
  
QY 1 WXXWXF 6  
| | |

Db 7 WTAWAF 12  
  
RESULT 3  
AAE26751  
ID AAE26751 standard; peptide; 9 AA.  
XX  
AC AAE26751;  
XX  
DT 13-DEC-2002 (first entry)  
XX  
DE Fibrin binding loop #3.  
XX  
KW Fibrin binding peptide; thrombosis; pulmonary embolism; atherosclerosis;  
KW myocardial infarct; ischaemia; imaging; rheumatoid arthritis; vasotropic;  
KW anaemia; hypoxia; tumour; diabetic retinopathy; autoimmune disorder;  
KW inflammatory disorder; angiogenesis; stroke; cerebroprotective.  
XX  
OS Unidentified.  
XX  
PN WO200255544-A2.  
XX  
PD 18-JUL-2002.  
XX  
PF 21-DEC-2001; 2001WO-US49534.  
XX  
PR 23-DEC-2000; 2000US-0747403.  
XX  
PA (DYAX-) DYAX CORP.  
XX  
PI Wescott CR, Beltzer JP, Sato AK;  
XX  
DR WPI; 2002-666875/71.  
XX  
PT Novel synthetic fibrin-binding moiety, useful for detecting, imaging or  
PT localizing fibrin-containing clots by magnetic resonance imaging,  
PT radioimaging and for treating diseases involving thrombus formation  
PT e.g. stroke -  
XX  
PS Claim 4; Page 55; 89pp; English.  
XX  
CC The invention relates to a synthetic fibrin binding group having affinity  
CC for fibrin. The invention is useful for detecting fibrin in a mammalian  
CC subject which involves (a) detectably labelling the binding group; (b)  
CC administering to the subject the labelled polypeptide, and (c) detecting  
CC the labelled polypeptide in the subject. The invention is useful for  
CC treating a disease involving thrombus formation eg. deep-vein thrombosis,  
CC pulmonary embolism, cardiogenic thrombosis, atherosclerosis, myocardial  
CC infarct, reperfusion ischaemia or stroke. The binding moieties are useful  
CC for detection, imaging and localisation of fibrin-containing clots by  
CC magnetic resonance imaging, radioimaging and other imaging methods and  
CC are also useful in the diagnosis and treatment of coronary conditions  
CC where fibrin plays a role. The fibrin binding moieties are useful for  
CC detecting and diagnosing numerous pathophysiologies in which fibrin plays  
CC a role eg. peritoneal adhesions which often occur after surgery or  
CC inflammatory and neoplastic processes and are comprised of a fibrin  
CC network, fibroblasts, macrophages and new blood vessels; rheumatoid  
CC arthritis, lupus or septic arthritis which often have bits of fibrin  
CC containing tissues called rice bodies in the synovial fluid of their  
CC joints; thrombocytopenic purpura, a type of anaemia in which deposits in  
CC arterioles causes turbulent blood flow resulting in stress and  
CC destruction of red blood cells. The fibrin specific agents can also be  
CC used to detect hypoxia or ischaemia of heart, kidney, liver, lung, brain  
CC or other organs, as well as the detection of tumours, diabetic  
CC retinopathy, early or high-risk atherosclerosis and other autoimmune and  
CC inflammatory disorders. Fibrin specific agents also could provide both  
CC direct or surrogate markers of disease models in which hypoxia and  
CC angiogenesis are expected to play a role. The invention is also useful  
CC for screening molecular libraries. The present sequence is a fibrin  
CC binding loop.  
XX  
SQ Sequence 9 AA;

Query Match 87.1%; Score 27; DB 23; Length 9;  
Best Local Similarity 50.0%; Pred. No. 9.3e+05;  
Matches 3; Conservative 0; Mismatches 3; Indels 0; Gaps 0;

QY 1 WXXWXF 6  
| | |  
Db 3 WESWTF 8

RESULT 4  
AAE26733  
ID AAE26733 standard; peptide; 15 AA.  
XX  
AC AAE26733;  
XX  
DT 13-DEC-2002 (first entry)  
XX  
DE Fibrin binding peptide #4.  
XX  
KW Fibrin binding peptide; thrombosis; pulmonary embolism; atherosclerosis;  
KW myocardial infarct; ischaemia; imaging; rheumatoid arthritis; vasotropic;  
KW anaemia; hypoxia; tumour; diabetic retinopathy; autoimmune disorder;  
KW inflammatory disorder; angiogenesis; stroke; cerebroprotective.  
XX  
OS Unidentified.  
XX  
PN WO200255544-A2.  
XX  
PD 18-JUL-2002.  
XX  
PF 21-DEC-2001; 2001WO-US49534.  
XX  
PR 23-DEC-2000; 2000US-0747403.  
XX  
PA (DYAX-) DYAX CORP.  
XX  
PI Wescott CR, Beltzer JP, Sato AK;  
XX  
DR WPI; 2002-666875/71.  
XX  
PT Novel synthetic fibrin-binding moiety, useful for detecting, imaging or  
PT localizing fibrin-containing clots by magnetic resonance imaging,  
PT radioimaging and for treating diseases involving thrombus formation  
PT e.g. stroke -  
XX  
PS Claim 10; Page 57; 89pp; English.  
XX  
CC The invention relates to a synthetic fibrin binding group having affinity  
CC for fibrin. The invention is useful for detecting fibrin in a mammalian  
CC subject which involves (a) detectably labelling the binding group; (b)  
CC administering to the subject the labelled polypeptide, and (c) detecting  
CC the labelled polypeptide in the subject. The invention is useful for  
CC treating a disease involving thrombus formation eg. deep-vein thrombosis,  
CC pulmonary embolism, cardiogenic thrombosis, atherosclerosis, myocardial  
CC infarct, reperfusion ischaemia or stroke. The binding moieties are useful  
CC for detection, imaging and localisation of fibrin-containing clots by  
CC magnetic resonance imaging, radioimaging and other imaging methods and  
CC are also useful in the diagnosis and treatment of coronary conditions  
CC where fibrin plays a role. The fibrin binding moieties are useful for  
CC detecting and diagnosing numerous pathophysiological in which fibrin plays  
CC a role eg. peritoneal adhesions which often occur after surgery or  
CC inflammatory and neoplastic processes and are comprised of a fibrin  
CC network, fibroblasts, macrophages and new blood vessels; rheumatoid  
CC arthritis, lupus or septic arthritis which often have bits of fibrin  
CC containing tissues called rice bodies in the synovial fluid of their  
CC joints; thrombocytopenic purpura, a type of anaemia in which deposits in  
CC arterioles causes turbulent blood flow resulting in stress and  
CC destruction of red blood cells. The fibrin specific agents can also be  
CC used to detect hypoxia or ischaemia of heart, kidney, liver, lung, brain  
CC or other organs, as well as the detection of tumours, diabetic  
CC retinopathy, early or high-risk atherosclerosis and other autoimmune and  
CC inflammatory disorders. Fibrin specific agents also could provide both  
CC direct or surrogate markers of disease models in which hypoxia and

CC angiogenesis are expected to play a role. The invention is also useful  
CC for screening molecular libraries. The present sequence is a fibrin  
CC binding peptide.

XX  
SQ Sequence 15 AA;  
Query Match 87.1%; Score 27; DB 23; Length 15;  
Best Local Similarity 50.0%; Pred. No. 1.6e+02;  
Matches 3; Conservative 0; Mismatches 3; Indels 0; Gaps 0;

QY 1 WXXWXF 6  
| | |  
Db 6 WESWTF 11

RESULT 5  
AAB01505  
ID AAB01505 standard; peptide; 6 AA.  
XX  
AC AAB01505;  
XX  
DT 08-NOV-2000 (first entry)  
XX  
DE Peptide which binds to transcription factor E2F-1 DNA binding domain.  
XX  
KW DNA binding; transcription factor; E2F; E2F-1; cell cycle; DP-1;  
KW activation; transcription; apoptosis; proliferative disorder;  
KW psoriasis; restenosis.  
XX  
OS Synthetic.  
XX  
PN WO200044771-A1.  
XX  
PD 03-AUG-2000.  
XX  
PF 26-JAN-2000; 2000WO-GB00227.  
XX  
PR 26-JAN-1999; 99GB-0001710.  
XX  
PA (PROL-) PROLIFIX LTD.  
XX  
PI Mueller R, Kontermann RE, Montigiani S;  
XX  
DR WPI; 2000-532806/48.  
XX  
PT Peptides binding to the DNA binding domain of transcription factor E2F  
PT and inhibiting cell cycle progression, useful for the treatment of  
PT cancer  
XX  
PS Example; Page 26; 42pp; English.  
XX  
CC Peptides which bind to the DNA binding domain of transcription  
CC factor E2F and inhibit cell cycle progression may be useful as  
CC research agents to investigate the interaction between E2F and DP-1,  
CC or the activation of transcription by E2F-1/DP-1 heterodimers. They  
CC may also be used for inducing apoptosis and/or cell cycle arrest in  
CC a cell, particularly for treatment of cancer or other proliferative  
CC disorders such as psoriasis and restenosis.  
XX  
SQ Sequence 6 AA;  
Query Match 83.9%; Score 26; DB 21; Length 6;  
Best Local Similarity 50.0%; Pred. No. 9.3e+05;  
Matches 3; Conservative 0; Mismatches 3; Indels 0; Gaps 0;

QY 1 WXXWXF 6  
| | |  
Db 1 WARWHF 6

RESULT 6  
AAB01506  
ID AAB01506 standard; peptide; 6 AA.

XX AAB01506;  
AC 08-NOV-2000 (first entry)  
XX  
XX  
DE Peptide which binds to transcription factor E2F-1 DNA binding domain.  
XX  
XX  
KW DNA binding; transcription factor; E2F; E2F-1; cell cycle; DP-1;  
KW activation; transcription; apoptosis; proliferative disorder;  
KW psoriasis; restenosis.  
XX  
OS Synthetic.  
XX  
PN WO200044771-A1.  
XX  
XX 03-AUG-2000.  
XX  
XX 26-JAN-2000; 2000WO-GB00227.  
XX  
PR 26-JAN-1999; 99GB-0001710.  
XX  
XX (PROL-) PROLIFIX LTD.  
XX  
PI Mueller R, Kontermann RE, Montigiani S;  
XX  
XX WPI; 2000-532806/48.  
XX  
XX Peptides binding to the DNA binding domain of transcription factor E2F  
and inhibiting cell cycle progression, useful for the treatment of  
cancer  
XX  
XX Example; Page 26; 42pp; English.  
XX  
XX Peptides which bind to the DNA binding domain of transcription  
factor E2F and inhibit cell cycle progression may be useful as  
research agents to investigate the interaction between E2F and DP-1,  
or the activation of transcription by E2F-1/DP-1 heterodimers. They  
may also be used for inducing apoptosis and/or cell cycle arrest in  
a cell, particularly for treatment of cancer or other proliferative  
disorders such as psoriasis and restenosis.  
XX  
SQ Sequence 6 AA;  
PS  
XX  
XX  
CC Peptides which bind to the DNA binding domain of transcription  
factor E2F and inhibit cell cycle progression may be useful as  
research agents to investigate the interaction between E2F and DP-1,  
or the activation of transcription by E2F-1/DP-1 heterodimers. They  
may also be used for inducing apoptosis and/or cell cycle arrest in  
a cell, particularly for treatment of cancer or other proliferative  
disorders such as psoriasis and restenosis.  
XX  
SQ Sequence 6 AA;  
PS  
XX  
XX  
CC Peptides which bind to the DNA binding domain of transcription  
factor E2F and inhibit cell cycle progression may be useful as  
research agents to investigate the interaction between E2F and DP-1,  
or the activation of transcription by E2F-1/DP-1 heterodimers. They  
may also be used for inducing apoptosis and/or cell cycle arrest in  
a cell, particularly for treatment of cancer or other proliferative  
disorders such as psoriasis and restenosis.  
XX  
SQ  
Query Match 83.9%; Score 26; DB 21; Length 6;  
Best Local Similarity 50.0%; Pred. No. 9.3e+05;  
Matches 3; Conservative 0; Mismatches 3; Indels 0; Gaps 0;  
QY 1 WXXWXF 6  
Db 1 WVAWHF 6  
RESULT 7  
AAB01508  
ID AAB01508 standard; peptide; 6 AA.  
XX  
AC AAB01508;  
XX  
XX 08-NOV-2000 (first entry)  
XX  
DE Peptide which binds to transcription factor E2F-1 DNA binding domain.  
XX  
XX  
KW DNA binding; transcription factor; E2F; E2F-1; cell cycle; DP-1;  
KW activation; transcription; apoptosis; proliferative disorder;  
KW psoriasis; restenosis.  
XX  
OS Synthetic.  
XX  
PN WO200044771-A1.  
XX  
XX 03-AUG-2000.  
XX  
XX 26-JAN-2000; 2000WO-GB00227.

XX 26-JAN-1999; 99GB-0001710.  
PR  
XX  
XX (PROL-) PROLIFIX LTD.  
XX  
XX  
PI Mueller R, Kontermann RE, Montigiani S;  
XX  
XX WPI; 2000-532806/48.  
XX  
XX Peptides binding to the DNA binding domain of transcription factor E2F  
and inhibiting cell cycle progression, useful for the treatment of  
cancer  
XX  
XX Example; Page 26; 42pp; English.  
XX  
XX Peptides which bind to the DNA binding domain of transcription  
factor E2F and inhibit cell cycle progression may be useful as  
research agents to investigate the interaction between E2F and DP-1,  
or the activation of transcription by E2F-1/DP-1 heterodimers. They  
may also be used for inducing apoptosis and/or cell cycle arrest in  
a cell, particularly for treatment of cancer or other proliferative  
disorders such as psoriasis and restenosis.  
XX  
SQ Sequence 6 AA;  
PS  
XX  
XX  
CC Peptides which bind to the DNA binding domain of transcription  
factor E2F and inhibit cell cycle progression may be useful as  
research agents to investigate the interaction between E2F and DP-1,  
or the activation of transcription by E2F-1/DP-1 heterodimers. They  
may also be used for inducing apoptosis and/or cell cycle arrest in  
a cell, particularly for treatment of cancer or other proliferative  
disorders such as psoriasis and restenosis.  
XX  
SQ  
Query Match 83.9%; Score 26; DB 21; Length 6;  
Best Local Similarity 50.0%; Pred. No. 9.3e+05;  
Matches 3; Conservative 0; Mismatches 3; Indels 0; Gaps 0;  
QY 1 WXXWXF 6  
Db 1 WVRWAF 6  
RESULT 8  
ABR45313  
ID ABR45313 standard; Peptide; 6 AA.  
XX  
AC ABR45313;  
XX  
XX 10-JUN-2003 (first entry)  
XX  
XX Staphylococcus aureus CHIPS-related peptide #503.  
DE  
XX  
XX CHIPS; Chemotaxis Inhibitory Protein; C5a-receptor; C5aR;  
formylated peptide receptor; FPR; neutrophil; monocyte; endothelial cell;  
inflammation; cardiovascular disease; central nervous system disease;  
gastrointestinal disease; skin disease; genitourinary disease;  
joint disease; respiratory disease; HIV infection; antiinflammatory;  
cardiant; cerebroprotective; neuroprotective; nootropic; dermatological;  
gynecological; immunosuppressive; anti-HIV.  
XX  
OS Staphylococcus aureus.  
OS Synthetic.  
XX  
XX WO2003006048-A1.  
XX  
XX 23-JAN-2003.  
XX  
XX 11-JUL-2001; 2001WO-EP08004.  
PF  
XX 11-JUL-2001; 2001WO-EP08004.  
PR  
XX (JARI-) JARI PHARM BV.  
PA  
XX Van Kessel CPM, Gosselaar-de Haas CJC, Kruijtzer JAW;  
PI Van Strijp JAG;  
XX  
XX WPI; 2003-247783/25.  
DR  
XX  
XX  
PT Combination of peptides derived from chemotaxis inhibiting protein from  
Staphylococcus aureus (CHIPS) having CHIPS activity, useful in  
prophylaxis and treatment of inflammation, cardiovascular, skin and

PT kidney diseases -  
 PS Disclosure; Page 12; 89pp; English.  
 XX  
 CC The present invention relates to peptides (ABR44811-ABR47162 and  
 CC ABR47164-ABR47385) derived from the Chemotaxis Inhibitory Protein (CHIPS)  
 CC from Staphylococcus aureus. The peptide fragments are useful in the  
 CC prophylaxis or treatment of diseases or disorders involving the  
 CC C5a-receptor (C5aR) and/or formylated peptide receptor (FPR) or  
 CC neutrophils, monocytes and endothelial cells or involving acute or  
 CC chronic inflammation reactions. The diseases or disorders include  
 CC cardiovascular diseases, disease of the central nervous system,  
 CC gastrointestinal diseases, skin diseases, genitourinary diseases, joint  
 CC diseases, respiratory diseases and HIV infection.  
 XX  
 SQ Sequence 6 AA;  
 Query Match 83.9%; Score 26; DB 24; Length 6;  
 Best Local Similarity 50.0%; Pred. No. 9.3e+05;  
 Matches 3; Conservative 0; Mismatches 3; Indels 0; Gaps 0;  
 QY 1 WXXWXF 6  
 | | |  
 Db 1 WSWFFF 6  
 RESULT 9  
 ABR45314  
 ID ABR45314 standard; Peptide; 6 AA.  
 AC ABR45314;  
 XX  
 DT 10-JUN-2003 (first entry)  
 XX  
 DE Staphylococcus aureus CHIPS-related peptide #504.  
 XX  
 KW CHIPS; Chemotaxis Inhibitory Protein; C5a-receptor; C5aR;  
 KW formylated peptide receptor; FPR; neutrophil; monocyte; endothelial cell;  
 KW inflammation; cardiovascular disease; central nervous system disease;  
 KW gastrointestinal disease; skin disease; genitourinary disease;  
 KW joint disease; respiratory disease; HIV infection; antiinflammatory;  
 KW cardiant; cerebroprotective; neuroprotective; nootropic; dermatological;  
 KW gynecological; immunosuppressive; anti-HIV.  
 XX  
 OS Staphylococcus aureus.  
 OS Synthetic.  
 OS  
 PN WO2003006048-A1.  
 XX  
 PD 23-JAN-2003.  
 XX  
 PF 11-JUL-2001; 2001WO-EP08004.  
 XX  
 PR 11-JUL-2001; 2001WO-EP08004.  
 XX  
 PA (JARI-) JARI PHARM BV.  
 XX  
 PI Van Kessel CPM, Gosselaar-de Haas CJC, Kruijtzer JAW;  
 PI Van Strijp JAG;  
 XX  
 DR WPI; 2003-247783/25.  
 XX  
 PT Combination of peptides derived from chemotaxis inhibiting protein from  
 PT Staphylococcus aureus (CHIPS) having CHIPS activity, useful in  
 PT prophylaxis and treatment of inflammation, cardiovascular, skin and  
 PT kidney diseases -  
 XX  
 PS Disclosure; Page 12; 89pp; English.  
 XX  
 CC The present invention relates to peptides (ABR44811-ABR47162 and  
 CC ABR47164-ABR47385) derived from the Chemotaxis Inhibitory Protein (CHIPS)  
 CC from Staphylococcus aureus. The peptide fragments are useful in the  
 CC prophylaxis or treatment of diseases or disorders involving the  
 CC C5a-receptor (C5aR) and/or formylated peptide receptor (FPR) or  
 CC neutrophils, monocytes and endothelial cells or involving acute or  
 CC chronic inflammation reactions. The diseases or disorders include  
 CC cardiovascular diseases, disease of the central nervous system,  
 CC gastrointestinal diseases, skin diseases, genitourinary diseases, joint  
 CC diseases, respiratory diseases and HIV infection.  
 XX  
 SQ Sequence 6 AA;

CC C5a-receptor (C5aR) and/or formylated peptide receptor (FPR) or  
 CC neutrophils, monocytes and endothelial cells or involving acute or  
 CC chronic inflammation reactions. The diseases or disorders include  
 CC cardiovascular diseases, disease of the central nervous system,  
 CC gastrointestinal diseases, skin diseases, genitourinary diseases, joint  
 CC diseases, respiratory diseases and HIV infection.  
 XX  
 SQ Sequence 6 AA;  
 Query Match 83.9%; Score 26; DB 24; Length 6;  
 Best Local Similarity 50.0%; Pred. No. 9.3e+05;  
 Matches 3; Conservative 0; Mismatches 3; Indels 0; Gaps 0;  
 QY 1 WXXWXF 6  
 | | |  
 Db 1 WTEWFF 6  
 RESULT 10  
 ABR45369  
 ID ABR45369 standard; Peptide; 6 AA.  
 XX  
 AC ABR45369;  
 XX  
 DT 10-JUN-2003 (first entry)  
 XX  
 DE Staphylococcus aureus CHIPS-related peptide #559.  
 XX  
 KW CHIPS; Chemotaxis Inhibitory Protein; C5a-receptor; C5aR;  
 KW formylated peptide receptor; FPR; neutrophil; monocyte; endothelial cell;  
 KW inflammation; cardiovascular disease; central nervous system disease;  
 KW gastrointestinal disease; skin disease; genitourinary disease;  
 KW joint disease; respiratory disease; HIV infection; antiinflammatory;  
 KW cardiant; cerebroprotective; neuroprotective; nootropic; dermatological;  
 KW gynecological; immunosuppressive; anti-HIV.  
 XX  
 OS Staphylococcus aureus.  
 OS Synthetic.  
 OS  
 PN WO2003006048-A1.  
 XX  
 PD 23-JAN-2003.  
 XX  
 PF 11-JUL-2001; 2001WO-EP08004.  
 XX  
 PR 11-JUL-2001; 2001WO-EP08004.  
 XX  
 PA (JARI-) JARI PHARM BV.  
 XX  
 PI Van Kessel CPM, Gosselaar-de Haas CJC, Kruijtzer JAW;  
 PI Van Strijp JAG;  
 XX  
 DR WPI; 2003-247783/25.  
 XX  
 PT Combination of peptides derived from chemotaxis inhibiting protein from  
 PT Staphylococcus aureus (CHIPS) having CHIPS activity, useful in  
 PT prophylaxis and treatment of inflammation, cardiovascular, skin and  
 PT kidney diseases -  
 XX  
 PS Disclosure; Page 12; 89pp; English.  
 XX  
 CC The present invention relates to peptides (ABR44811-ABR47162 and  
 CC ABR47164-ABR47385) derived from the Chemotaxis Inhibitory Protein (CHIPS)  
 CC from Staphylococcus aureus. The peptide fragments are useful in the  
 CC prophylaxis or treatment of diseases or disorders involving the  
 CC C5a-receptor (C5aR) and/or formylated peptide receptor (FPR) or  
 CC neutrophils, monocytes and endothelial cells or involving acute or  
 CC chronic inflammation reactions. The diseases or disorders include  
 CC cardiovascular diseases, disease of the central nervous system,  
 CC gastrointestinal diseases, skin diseases, genitourinary diseases, joint  
 CC diseases, respiratory diseases and HIV infection.  
 XX  
 SQ Sequence 6 AA;



Query Match 83.9%; Score 26; DB 24; Length 6;  
Best Local Similarity 50.0%; Pred. No. 9.3e+05;  
Matches 3; Conservative 0; Mismatches 3; Indels 0; Gaps 0;

QY 1 WXXWXF 6  
| | | |  
Db 1 WSWWIF 6

RESULT 11  
ABR45370  
ID ABR45370 standard; Peptide; 6 AA.  
XX  
AC ABR45370;  
XX  
DT 10-JUN-2003 (first entry)  
XX  
DE Staphylococcus aureus CHIPS-related peptide #560.  
XX  
KW CHIPS; Chemotaxis Inhibitory Protein; C5a-receptor; C5aR;  
KW formylated peptide receptor; FPR; neutrophil; monocyte; endothelial cell;  
KW inflammation; cardiovascular disease; central nervous system disease;  
KW gastrointestinal disease; skin disease; genitourinary disease;  
KW joint disease; respiratory disease; HIV infection; antiinflammatory;  
KW cardiant; cerebroprotective; neuroprotective; nootropic; dermatological;  
KW gynecological; immunosuppressive; anti-HIV.  
XX  
OS Staphylococcus aureus.  
OS Synthetic.  
XX  
PN WO2003006048-A1.  
XX  
PD 23-JAN-2003.  
XX  
PF 11-JUL-2001; 2001WO-EP08004.  
XX  
PR 11-JUL-2001; 2001WO-EP08004.  
XX  
PA (JARI-) JARI PHARM BV.  
XX  
PI Van Kessel CPM, Gosselaar-de Haas CJC, Kruijtzer JAW;  
PI Van Strijp JAG;  
XX  
WPI; 2003-247783/25.  
XX  
XX  
PT Combination of peptides derived from chemotaxis inhibiting protein from  
PT Staphylococcus aureus (CHIPS) having CHIPS activity, useful in  
PT prophylaxis or treatment of inflammation, cardiovascular, skin and  
PT kidney diseases -  
XX  
PS Disclosure; Page 12; 89pp; English.  
XX  
CC The present invention relates to peptides (ABR44811-ABR47162 and  
CC ABR47164-ABR47385) derived from the Chemotaxis Inhibitory Protein (CHIPS)  
CC from Staphylococcus aureus. The peptide fragments are useful in the  
CC prophylaxis or treatment of diseases or disorders involving the  
CC C5a-receptor (C5aR) and/or formylated peptide receptor (FPR) or  
CC neutrophils, monocytes and endothelial cells or involving acute or  
CC chronic inflammation reactions. The diseases or disorders include  
CC cardiovascular diseases, disease of the central nervous system,  
CC gastrointestinal diseases, skin diseases, genitourinary diseases, joint  
XX diseases, respiratory diseases and HIV infection.

Sequence 6 AA;  
Query Match 83.9%; Score 26; DB 24; Length 6;  
Best Local Similarity 50.0%; Pred. No. 9.3e+05;  
Matches 3; Conservative 0; Mismatches 3; Indels 0; Gaps 0;

QY 1 WXXWXF 6  
| | | |  
Db 1 WTFWIF 6

RESULT 12  
ABR45425  
ID ABR45425 standard; Peptide; 6 AA.  
XX  
AC ABR45425;  
XX  
DT 10-JUN-2003 (first entry)  
XX  
DE Staphylococcus aureus CHIPS-related peptide #615.  
XX  
KW CHIPS; Chemotaxis Inhibitory Protein; C5a-receptor; C5aR;  
KW formylated peptide receptor; FPR; neutrophil; monocyte; endothelial cell;  
KW inflammation; cardiovascular disease; central nervous system disease;  
KW gastrointestinal disease; skin disease; genitourinary disease;  
KW joint disease; respiratory disease; HIV infection; antiinflammatory;  
KW cardiant; cerebroprotective; neuroprotective; nootropic; dermatological;  
KW gynecological; immunosuppressive; anti-HIV.  
XX  
OS Staphylococcus aureus.  
OS Synthetic.  
XX  
PN WO2003006048-A1.  
XX  
PD 23-JAN-2003.  
XX  
PF 11-JUL-2001; 2001WO-EP08004.  
XX  
PR 11-JUL-2001; 2001WO-EP08004.  
XX  
PA (JARI-) JARI PHARM BV.  
XX  
PI Van Kessel CPM, Gosselaar-de Haas CJC, Kruijtzer JAW;  
PI Van Strijp JAG;  
XX  
WPI; 2003-247783/25.  
XX  
XX  
PT Combination of peptides derived from chemotaxis inhibiting protein from  
PT Staphylococcus aureus (CHIPS) having CHIPS activity, useful in  
PT prophylaxis and treatment of inflammation, cardiovascular, skin and  
PT kidney diseases -  
XX  
PS Disclosure; Page 12; 89pp; English.  
XX  
CC The present invention relates to peptides (ABR44811-ABR47162 and  
CC ABR47164-ABR47385) derived from the Chemotaxis Inhibitory Protein (CHIPS)  
CC from Staphylococcus aureus. The peptide fragments are useful in the  
CC prophylaxis or treatment of diseases or disorders involving the  
CC C5a-receptor (C5aR) and/or formylated peptide receptor (FPR) or  
CC neutrophils, monocytes and endothelial cells or involving acute or  
CC chronic inflammation reactions. The diseases or disorders include  
CC cardiovascular diseases, disease of the central nervous system,  
CC gastrointestinal diseases, skin diseases, genitourinary diseases, joint  
XX diseases, respiratory diseases and HIV infection.

Sequence 6 AA;  
Query Match 83.9%; Score 26; DB 24; Length 6;  
Best Local Similarity 50.0%; Pred. No. 9.3e+05;  
Matches 3; Conservative 0; Mismatches 3; Indels 0; Gaps 0;

QY 1 WXXWXF 6  
| | | |  
Db 1 WSWWLF 6

RESULT 13  
ABR45426  
ID ABR45426 standard; Peptide; 6 AA.  
XX  
AC ABR45426;  
XX



DT 10-JUN-2003 (first entry)  
XX  
DE Staphylococcus aureus CHIPS-related peptide #616.  
XX  
KW CHIPS; Chemotaxis Inhibitory Protein; C5a-receptor; C5aR;  
KW formylated peptide receptor; FPR; neutrophil; monocyte; endothelial cell;  
KW inflammation; cardiovascular disease; central nervous system disease;  
KW gastrointestinal disease; skin disease; genitourinary disease;  
KW joint disease; respiratory disease; HIV infection; antiinflammatory;  
KW cardiant; cerebroprotective; neuroprotective; nootropic; dermatological;  
KW gynecological; immunosuppressive; anti-HIV.  
XX  
OS Staphylococcus aureus.  
OS Synthetic.  
XX  
PN WO2003006048-A1.  
XX  
PD 23-JAN-2003.  
XX  
PF 11-JUL-2001; 2001WO-EP08004.  
XX  
PR 11-JUL-2001; 2001WO-EP08004.  
XX  
PA (JARI-) JARI PHARM BV.  
XX  
PI Van Kessel CPM, Gosselaar-de Haas CJC, Kruijtzer JAW;  
PI Van Strijp JAG;  
XX  
DR WPI; 2003-247783/25.  
XX  
PT Combination of peptides derived from chemotaxis inhibiting protein from  
PT Staphylococcus aureus (CHIPS) having CHIPS activity, useful in  
PT prophylaxis and treatment of inflammation, cardiovascular, skin and  
PT kidney diseases -  
XX  
PS Disclosure; Page 12; 89pp; English.  
XX  
CC The present invention relates to peptides (ABR44811-ABR47162 and  
CC ABR47164-ABR47385) derived from the Chemotaxis Inhibitory Protein (CHIPS)  
CC from Staphylococcus aureus. The peptide fragments are useful in the  
CC prophylaxis or treatment of diseases or disorders involving the  
CC C5a-receptor (C5aR) and/or formylated peptide receptor (FPR) or  
CC neutrophils, monocytes and endothelial cells or involving acute or  
CC chronic inflammation reactions. The diseases or disorders include  
CC cardiovascular diseases, disease of the central nervous system,  
CC gastrointestinal diseases, skin diseases, genitourinary diseases, joint  
CC diseases, respiratory diseases and HIV infection.  
XX  
SQ Sequence 6 AA;  
  
Query Match 83.9%; Score 26; DB 24; Length 6;  
Best Local Similarity 50.0%; Pred. No. 9.3e+05;  
Matches 3; Conservative 0; Mismatches 3; Indels 0; Gaps 0;  
  
QY 1 WXXWXF 6  
| | | |  
Db 1 WTFWLF 6  
  
RESULT 14  
ABR45481  
ID ABR45481 standard; Peptide; 6 AA.  
XX  
AC ABR45481;  
XX  
DT 10-JUN-2003 (first entry)  
XX  
DE Staphylococcus aureus CHIPS-related peptide #671.  
XX  
KW CHIPS; Chemotaxis Inhibitory Protein; C5a-receptor; C5aR;  
KW formylated peptide receptor; FPR; neutrophil; monocyte; endothelial cell;  
KW inflammation; cardiovascular disease; central nervous system disease;  
KW gastrointestinal disease; skin disease; genitourinary disease;

KW joint disease; respiratory disease; HIV infection; antiinflammatory;  
KW cardiant; cerebroprotective; neuroprotective; nootropic; dermatological;  
KW gynecological; immunosuppressive; anti-HIV.  
XX  
OS Staphylococcus aureus.  
OS Synthetic.  
XX  
PN WO2003006048-A1.  
XX  
PD 23-JAN-2003.  
XX  
PF 11-JUL-2001; 2001WO-EP08004.  
XX  
PR 11-JUL-2001; 2001WO-EP08004.  
XX  
PA (JARI-) JARI PHARM BV.  
XX  
PI Van Kessel CPM, Gosselaar-de Haas CJC, Kruijtzer JAW;  
PI Van Strijp JAG;  
XX  
DR WPI; 2003-247783/25.  
XX  
PT Combination of peptides derived from chemotaxis inhibiting protein from  
PT Staphylococcus aureus (CHIPS) having CHIPS activity, useful in  
PT prophylaxis and treatment of inflammation, cardiovascular, skin and  
PT kidney diseases -  
XX  
PS Disclosure; Page 13; 89pp; English.  
XX  
CC The present invention relates to peptides (ABR44811-ABR47162 and  
CC ABR47164-ABR47385) derived from the Chemotaxis Inhibitory Protein (CHIPS)  
CC from Staphylococcus aureus. The peptide fragments are useful in the  
CC prophylaxis or treatment of diseases or disorders involving the  
CC C5a-receptor (C5aR) and/or formylated peptide receptor (FPR) or  
CC neutrophils, monocytes and endothelial cells or involving acute or  
CC chronic inflammation reactions. The diseases or disorders include  
CC cardiovascular diseases, disease of the central nervous system,  
CC gastrointestinal diseases, skin diseases, genitourinary diseases, joint  
CC diseases, respiratory diseases and HIV infection.  
XX  
SQ Sequence 6 AA;  
  
Query Match 83.9%; Score 26; DB 24; Length 6;  
Best Local Similarity 50.0%; Pred. No. 9.3e+05;  
Matches 3; Conservative 0; Mismatches 3; Indels 0; Gaps 0;  
  
QY 1 WXXWXF 6  
| | | |  
Db 1 WSPWVF 6  
  
RESULT 15  
ABR45482  
ID ABR45482 standard; Peptide; 6 AA.  
XX  
AC ABR45482;  
XX  
DT 10-JUN-2003 (first entry)  
XX  
DE Staphylococcus aureus CHIPS-related peptide #672.  
XX  
KW CHIPS; Chemotaxis Inhibitory Protein; C5a-receptor; C5aR;  
KW formylated peptide receptor; FPR; neutrophil; monocyte; endothelial cell;  
KW inflammation; cardiovascular disease; central nervous system disease;  
KW gastrointestinal disease; skin disease; genitourinary disease;  
KW joint disease; respiratory disease; HIV infection; antiinflammatory;  
KW cardiant; cerebroprotective; neuroprotective; nootropic; dermatological;  
KW gynecological; immunosuppressive; anti-HIV.  
XX  
OS Staphylococcus aureus.  
OS Synthetic.  
XX  
PN WO2003006048-A1.

XX 23-JAN-2003.  
PD  
XX  
PF 11-JUL-2001; 2001WO-EP08004.  
XX  
PR 11-JUL-2001; 2001WO-EP08004.  
XX  
PA (JARI-) JARI PHARM BV.  
XX  
PI Van Kessel CPM, Gosselaar-de Haas CJC, Kruijtzer JAW;  
PI Van Strijp JAG;  
XX  
DR WPI; 2003-247783/25.  
XX  
XX Combination of peptides derived from chemotaxis inhibiting protein from  
PT Staphylococcus aureus (CHIPS) having CHIPS activity, useful in  
PT prophylaxis and treatment of inflammation, cardiovascular, skin and  
PT kidney diseases -  
XX  
PS Disclosure; Page 13; 89pp; English.  
XX  
CC The present invention relates to peptides (ABR44811-ABR47162 and  
CC ABR47164-ABR47385) derived from the Chemotaxis Inhibitory Protein (CHIPS)  
CC from Staphylococcus aureus. The peptide fragments are useful in the  
CC prophylaxis or treatment of diseases or disorders involving the  
CC C5a-receptor (C5aR) and/or formylated peptide receptor (FPR) or  
CC neutrophils, monocytes and endothelial cells or involving acute or  
CC chronic inflammation reactions. The diseases or disorders include  
CC cardiovascular diseases, disease of the central nervous system,  
CC gastrointestinal diseases, skin diseases, genitourinary diseases, joint  
CC diseases, respiratory diseases and HIV infection.  
XX  
SQ Sequence 6 AA;  
  
Query Match 83.9%; Score 26; DB 24; Length 6;  
Best Local Similarity 50.0%; Pred. No. 9.3e+05;  
Matches 3; Conservative 0; Mismatches 3; Indels 0; Gaps 0;  
  
QY 1 WXXWXF 6  
| | |  
Db 1 WTFWVF 6  
  
RESULT 16  
ABR45593  
ID ABR45593 standard; Peptide; 6 AA.  
XX  
AC ABR45593;  
XX  
DT 10-JUN-2003 (first entry)  
XX  
DE Staphylococcus aureus CHIPS-related peptide #783.  
XX  
KW CHIPS; Chemotaxis Inhibitory Protein; C5a-receptor; C5aR;  
KW formylated peptide receptor; FPR; neutrophil; monocyte; endothelial cell;  
KW inflammation; cardiovascular disease; central nervous system disease;  
KW gastrointestinal disease; skin disease; genitourinary disease;  
KW joint disease; respiratory disease; HIV infection; antiinflammatory;  
KW cardiant; cerebroprotective; neuroprotective; nootropic; dermatological;  
KW gynecological; immunosuppressive; anti-HIV.  
XX  
OS Staphylococcus aureus.  
OS Synthetic.  
XX  
PN WO2003006048-A1.  
XX  
PD 23-JAN-2003.  
XX  
PF 11-JUL-2001; 2001WO-EP08004.  
XX  
PR 11-JUL-2001; 2001WO-EP08004.  
XX  
PA (JARI-) JARI PHARM BV.

XX Van Kessel CPM, Gosselaar-de Haas CJC, Kruijtzer JAW;  
PI Van Strijp JAG;  
XX  
DR WPI; 2003-247783/25.  
XX  
XX Combination of peptides derived from chemotaxis inhibiting protein from  
PT Staphylococcus aureus (CHIPS) having CHIPS activity, useful in  
PT prophylaxis and treatment of inflammation, cardiovascular, skin and  
PT kidney diseases -  
XX  
PS Disclosure; Page 13; 89pp; English.  
XX  
CC The present invention relates to peptides (ABR44811-ABR47162 and  
CC ABR47164-ABR47385) derived from the Chemotaxis Inhibitory Protein (CHIPS)  
CC from Staphylococcus aureus. The peptide fragments are useful in the  
CC prophylaxis or treatment of diseases or disorders involving the  
CC C5a-receptor (C5aR) and/or formylated peptide receptor (FPR) or  
CC neutrophils, monocytes and endothelial cells or involving acute or  
CC chronic inflammation reactions. The diseases or disorders include  
CC cardiovascular diseases, disease of the central nervous system,  
CC gastrointestinal diseases, skin diseases, genitourinary diseases, joint  
CC diseases, respiratory diseases and HIV infection.  
XX  
SQ Sequence 6 AA;  
  
Query Match 83.9%; Score 26; DB 24; Length 6;  
Best Local Similarity 50.0%; Pred. No. 9.3e+05;  
Matches 3; Conservative 0; Mismatches 3; Indels 0; Gaps 0;  
  
QY 1 WXXWXF 6  
| | |  
Db 1 WSWFYF 6  
  
RESULT 17  
ABR45594  
ID ABR45594 standard; Peptide; 6 AA.  
XX  
AC ABR45594;  
XX  
DT 10-JUN-2003 (first entry)  
XX  
DE Staphylococcus aureus CHIPS-related peptide #784.  
XX  
KW CHIPS; Chemotaxis Inhibitory Protein; C5a-receptor; C5aR;  
KW formylated peptide receptor; FPR; neutrophil; monocyte; endothelial cell;  
KW inflammation; cardiovascular disease; central nervous system disease;  
KW gastrointestinal disease; skin disease; genitourinary disease;  
KW joint disease; respiratory disease; HIV infection; antiinflammatory;  
KW cardiant; cerebroprotective; neuroprotective; nootropic; dermatological;  
KW gynecological; immunosuppressive; anti-HIV.  
XX  
OS Staphylococcus aureus.  
OS Synthetic.  
XX  
PN WO2003006048-A1.  
XX  
PD 23-JAN-2003.  
XX  
PF 11-JUL-2001; 2001WO-EP08004.  
XX  
PR 11-JUL-2001; 2001WO-EP08004.  
XX  
PA (JARI-) JARI PHARM BV.  
XX  
PI Van Kessel CPM, Gosselaar-de Haas CJC, Kruijtzer JAW;  
PI Van Strijp JAG;  
XX  
DR WPI; 2003-247783/25.  
XX  
PT Combination of peptides derived from chemotaxis inhibiting protein from  
PT Staphylococcus aureus (CHIPS) having CHIPS activity, useful in

PT prophylaxis and treatment of inflammation, cardiovascular, skin and  
PT kidney diseases -  
PS Disclosure; Page 13; 89pp; English.  
XX  
CC The present invention relates to peptides (ABR44811-ABR47162 and  
CC ABR47164-ABR47385) derived from the Chemotaxis Inhibitory Protein (CHIPS)  
CC from Staphylococcus aureus. The peptide fragments are useful in the  
CC prophylaxis or treatment of diseases or disorders involving the  
CC C5a-receptor (C5aR) and/or formulated peptide receptor (PPR) or  
CC neutrophils, monocytes and endothelial cells or involving acute or  
CC chronic inflammation reactions. The diseases or disorders include  
CC cardiovascular diseases, disease of the central nervous system,  
CC gastrointestinal diseases, skin diseases, genitourinary diseases, joint  
CC diseases, respiratory diseases and HIV infection.  
XX  
SQ Sequence 6 AA;  
Query Match 83.9%; Score 26; DB 24; Length 6;  
Best Local Similarity 50.0%; Pred. No. 9.3e+05;  
Matches 3; Conservative 0; Mismatches 3; Indels 0; Gaps 0;  
QY 1 WXXWXF 6  
Db 1 WTFWYF 6  
RESULT 18  
AAE26775  
ID AAE26775 standard; peptide; 9 AA.  
XX  
AC AAE26775;  
XX  
DT 13-DEC-2002 (first entry)  
DE  
DE Fibrin binding peptide #28.  
XX  
KW Fibrin binding peptide; thrombosis; pulmonary embolism; atherosclerosis;  
KW myocardial infarct; ischaemia; imaging; rheumatoid arthritis; vasotropic;  
KW anaemia; hypoxia; tumour; diabetic retinopathy; autoimmune disorder;  
KW inflammatory disorder; angiogenesis; stroke; cerebroprotective.  
XX  
OS Unidentified.  
XX  
XX WO200255544-A2.  
PN  
XX  
PD 18-JUL-2002.  
XX  
XX 21-DEC-2001; 2001WO-US49534.  
PF  
XX  
XX 23-DEC-2000; 2000US-0747403.  
PR  
XX  
PA (DYAX-) DYAX CORP.  
XX  
PI Wescott CR, Beltzer JP, Sato AK;  
XX  
XX WPI; 2002-666875/71.  
XX  
XX Novel synthetic fibrin-binding moiety, useful for detecting, imaging or  
PT localizing fibrin-containing clots by magnetic resonance imaging,  
PT radioimaging and for treating diseases involving thrombus formation  
PT e.g. stroke -  
XX  
XX Claim 4; Page 55; 89pp; English.  
PS  
XX  
CC The invention relates to a synthetic fibrin binding group having affinity  
CC for fibrin. The invention is useful for detecting fibrin in a mammalian  
CC subject which involves (a) detectably labelling the binding group; (b)  
CC administering to the subject the labelled polypeptide, and (c) detecting  
CC the labelled polypeptide in the subject. The invention is useful for  
CC treating a disease involving thrombus formation eg. deep-vein thrombosis,  
CC pulmonary embolism, cardiogenic thrombosis, atherosclerosis, myocardial  
CC infarct, reperfusion ischaemia or stroke. The binding moieties are useful

CC for detection, imaging and localisation of fibrin-containing clots by  
CC magnetic resonance imaging, radioimaging and other imaging methods and  
CC are also useful in the diagnosis and treatment of coronary conditions  
CC where fibrin plays a role. The fibrin binding moieties are useful for  
CC detecting and diagnosing numerous pathophysiological in which fibrin plays  
CC a role eg. peritoneal adhesions which often occur after surgery or  
CC inflammatory and neoplastic processes and are comprised of a fibrin  
CC network, fibroblasts, macrophages and new blood vessels; rheumatoid  
CC arthritis, lupus or septic arthritis which often have bits of fibrin  
CC containing tissues called rice bodies in the synovial fluid of their  
CC joints; thrombocytopenic purpura, a type of anaemia in which deposits in  
CC arterioles causes turbulent blood flow resulting in stress and  
CC destruction of red blood cells. The fibrin specific agents can also be  
CC used to detect hypoxia or ischaemia of heart, kidney, liver, lung, brain  
CC or other organs, as well as the detection of tumours, diabetic  
CC retinopathy, early or high-risk atherosclerosis and other autoimmune and  
CC inflammatory disorders. Fibrin specific agents also could provide both  
CC direct or surrogate markers of disease models in which hypoxia and  
CC angiogenesis are expected to play a role. The invention is also useful  
CC for screening molecular libraries. The present sequence is a fibrin  
CC binding peptide.  
XX  
SQ Sequence 9 AA;  
Query Match 83.9%; Score 26; DB 23; Length 9;  
Best Local Similarity 50.0%; Pred. No. 9.3e+05;  
Matches 3; Conservative 0; Mismatches 3; Indels 0; Gaps 0;  
QY 1 WXXWXF 6  
Db 3 WGSWKF 8  
RESULT 19  
AAE65508  
ID AAY65508 standard; Peptide; 15 AA.  
XX  
AC AAY65508;  
XX  
DT 01-FEB-2000 (first entry)  
XX  
DE Oestrogen receptor alpha binding peptide 5PT.  
XX  
KW Oestrogen receptor; estrogen; estradiol; oestrogen response element;  
KW ERE; binding; biological activity; fingerprint; molecular braille;  
KW cellular braille; modulation; tamoxifen; breast cancer; ovarian cancer;  
KW menopause; osteoporosis; selective oestrogen receptor modulator;  
KW identification; characterisation; classification.  
XX  
OS Synthetic.  
OS Homo sapiens.  
XX  
PN WO9954728-A2.  
XX  
PD 28-OCT-1999.  
XX  
PF 26-MAR-1999; 99WO-US06664.  
XX  
XX 23-APR-1998; 98US-0082756.  
PR 09-SEP-1998; 98US-0099656.  
PR 08-JAN-1999; 99US-0115345.  
XX  
PA (NOVA-) NOVALON PHARM CORP.  
XX  
XX Paige LA, Hamilton PT, Fowlkes DM, Buehrer B, Barnett T;  
PI McDonnell DP, Christensen DJ;  
XX  
XX WPI; 2000-013281/01.  
XX  
PT Methods for identifying new receptor modulators, especially estrogen  
PT modulators to treat tamoxifen refractory breast cancer -  
XX  
PS Example 2.1; Page 159; 219pp; English.

XX The present invention describes a method for predicting the biological  
CC activity of new receptor modulating compounds (I) using novel oligomeric  
CC peptides (biokeys) which have differential abilities to bind to 2  
CC different receptor conformations. The method is used to identify new  
CC drugs that are physiological or pharmacological agonists/antagonists and  
CC that target various receptors, which are involved in certain disease  
CC conditions. The system may be used as a primary screening tool to  
CC identify hits, to classify lead compounds from a drug screen to,  
CC characterise selective oestrogen receptor modulators (SERMs) in terms of  
CC agonist and antagonist function and to predict possible clinical effects  
CC of SERMs such as tissue and receptor specificity. The method can also be  
CC applied to the fractionation of mixtures of SERMs to determine which  
CC components are producing agonistic and antagonistic activity. The method  
CC may be used with other receptors (e.g. progesterone, androgen,  
CC glucocorticoid, thyroid, vitamin D, beta-adrenergic, dopamine and  
CC epidermal growth factor, to identify, characterise and classify  
CC modulators of receptor activity. Peptides comprising a LXXLL motif may  
CC be used to modulate the oestrogen receptor in treating e.g. breast and  
CC ovarian cancer and ameliorating the effects of menopause, including  
CC osteoporosis. AAY65439 to AAY65652 represent oestrogen receptor,  
CC estradiol receptor and oestrogen response element binding peptides given  
CC in the exemplification of the present invention. AAZ35740 to AAZ35745  
CC represent oligonucleotides used in the exemplification of the present  
CC invention.

XX Sequence 15 AA;  
SQ

Query Match 83.9%; Score 26; DB 21; Length 15;  
Best Local Similarity 50.0%; Pred. No. 2.4e+02;  
Matches 3; Conservative 0; Mismatches 3; Indels 0; Gaps 0;

QY 1 WXXWXP 6  
| | |  
Db 8 WYDWTF 13

RESULT 20  
ABB99042  
ID ABB99042 standard; Peptide; 15 AA.  
XX  
AC ABB99042;  
XX  
DT 24-JAN-2003 (first entry)  
XX  
DE Serine/threonine protein kinase 9.13 N-terminal peptide sequence.  
XX  
KW Serine/threonine protein kinase 9.13; enzyme; tumour;  
KW embryonic development malformation; protein metabolic disorder.  
XX  
OS Unidentified.  
XX  
FN CN1352273-A.  
XX  
PD 05-JUN-2002.  
XX  
PF 02-NOV-2000; 2000CN-0127173.  
XX  
PR 02-NOV-2000; 2000CN-0127173.  
XX  
PA (BODE-) BODE GENE DEV CO LTD SHANGHAI.  
XX  
PI Mao Y, Xie Y;  
XX  
DR WPI; 2002-637138/69.  
XX  
PT New serine/threonine protein kinase 9.13 polypeptide for treating  
PT embryonic development malformation, various tumours and protein  
PT metabolic disorder -  
XX  
PS Example 5; Page 21 (disclosure); 35pp; Chinese.  
XX  
CC The present invention discloses a serine/threonine protein kinase 9.13,

CC the polynucleotides encoding the polypeptide, and a DNA recombination  
CC process to produce the polypeptide. The present invention also discloses  
CC applying the polypeptide in treating various diseases, such as embryonic  
CC development malformation, various tumours and protein metabolic disorder.  
CC The present invention also discloses the antagonist resisting the  
CC polypeptide and its treatment effect. The current sequence represents the  
CC serine/threonine protein kinase 9.13 N-terminal peptide sequence.

XX Sequence 15 AA;  
SQ

Query Match 83.9%; Score 26; DB 23; Length 15;  
Best Local Similarity 50.0%; Pred. No. 2.4e+02;  
Matches 3; Conservative 0; Mismatches 3; Indels 0; Gaps 0;

QY 1 WXXWXP 6  
| | |  
Db 6 WLFWSP 11

RESULT 21  
AAE26759  
ID AAE26759 standard; peptide; 15 AA.  
XX  
AC AAE26759;  
XX  
DT 13-DEC-2002 (first entry)  
XX  
DE Fibrin binding peptide #12.  
XX  
KW Fibrin binding peptide; thrombosis; pulmonary embolism; atherosclerosis;  
KW myocardial infarct; ischaemia; imaging; rheumatoid arthritis; vasotropic;  
KW anaemia; hypoxia; tumour; diabetic retinopathy; autoimmune disorder;  
KW inflammatory disorder; angiogenesis; stroke; cerebroprotective.  
XX  
OS Unidentified.  
XX  
FN WO200255544-A2.  
XX  
PD 18-JUL-2002.  
XX  
PF 21-DEC-2001; 2001WO-US49534.  
XX  
PR 23-DEC-2000; 2000US-0747403.  
XX  
PA (DYAX-) DYAX CORP.  
XX  
PI Wescott CR, Beltzer JP, Sato AK;  
XX  
DR WPI; 2002-666875/71.  
XX  
PT Novel synthetic fibrin-binding moiety, useful for detecting, imaging or  
PT localizing fibrin-containing clots by magnetic resonance imaging,  
PT radioimaging and for treating diseases involving thrombus formation  
PT e.g. stroke -  
XX  
PS Claim 10; Page 58; 89pp; English.  
XX  
CC The invention relates to a synthetic fibrin binding group having affinity  
CC for fibrin. The invention is useful for detecting fibrin in a mammalian  
CC subject which involves (a) detectably labelling the binding group; (b)  
CC administering to the subject the labelled polypeptide, and (c) detecting  
CC the labelled polypeptide in the subject. The invention is useful for  
CC treating a disease involving thrombus formation eg. deep-vein thrombosis,  
CC pulmonary embolism, cardiogenic thrombosis, atherosclerosis, myocardial  
CC infarct, reperfusion ischaemia or stroke. The binding moieties are useful  
CC for detection, imaging and localisation of fibrin-containing clots by  
CC magnetic resonance imaging, radioimaging and other imaging methods and  
CC are also useful in the diagnosis and treatment of coronary conditions  
CC where fibrin plays a role. The fibrin binding moieties are useful for  
CC detecting and diagnosing numerous pathophysiologies in which fibrin plays  
CC a role eg. peritoneal adhesions which often occur after surgery or  
CC inflammatory and neoplastic processes and are comprised of a fibrin  
CC network, fibroblasts, macrophages and new blood vessels; rheumatoid



CC arthritis, lupus or septic arthritis which often have bits of fibrin  
CC containing tissues called rice bodies in the synovial fluid of their  
CC joints; thrombocytopenic purpura, a type of anaemia in which deposits in  
CC arterioles causes turbulent blood flow resulting in stress and  
CC destruction of red blood cells. The fibrin specific agents can also be  
CC used to detect hypoxia or ischaemia of heart, kidney, liver, lung, brain  
CC or other organs, as well as the detection of tumours, diabetic  
CC retinopathy, early or high-risk atherosclerosis and other autoimmune and  
CC inflammatory disorders. Fibrin specific agents also could provide both  
CC direct or surrogate markers of disease models in which hypoxia and  
CC angiogenesis are expected to play a role. The invention is also useful  
CC for screening molecular libraries. The present sequence is a fibrin  
XX binding peptide.  
SQ Sequence 15 AA;  
Query Match 83.9%; Score 26; DB 23; Length 15;  
Best Local Similarity 50.0%; Pred. No. 2.4e+02;  
Matches 3; Conservative 0; Mismatches 3; Indels 0; Gaps 0;  
QY 1 WXXWXP 6  
Db 6 WGSWKF 11  
RESULT 22  
AAU86245  
ID AAU86245 standard; Peptide; 15 AA.  
XX  
AC AAU86245;  
XX  
DT 21-MAY-2002 (first entry)  
XX  
DE Oestrogen receptor alpha binding peptide 5PT.  
XX  
KW Oestrogen receptor; breast cancer; combinatorial peptide library;  
KW receptor modulating compound.  
XX Synthetic.  
OS  
XX WO200204956-A2.  
PN  
XX  
PD 17-JAN-2002.  
XX  
PF 11-JUL-2001; 2001WO-US21867.  
XX  
PR 12-JUL-2000; 2000US-0614865.  
PR 21-MAY-2001; 2001US-0860688.  
XX  
PA (KARO-) KARO BIO USA INC.  
XX  
PI Fowlkes DM, Barnett TR, Buehrer B;  
XX  
DR WPI; 2002-154969/20.  
XX  
XX Identifying receptor-binding peptides comprises screening combinatorial  
PT peptide library presented in form of cells each of which coexpress one  
PT peptide member and receptor with signal producing system for reporting  
PT binding -  
XX  
PS Disclosure; Page 142; 175pp; English.  
XX  
CC The invention relates to identifying a binding peptide which binds a  
CC receptor and which is a member of a combinatorial library of peptides,  
CC comprising screening a combinatorial peptide library presented in the  
CC form of cells which coexpress the receptor or its ligand-binding receptor  
CC moiety and one member of the library, together with a signal producing  
CC system for reporting binding of the peptide to the receptor. Also  
CC included is a method for predicting the receptor-modulating activity of a  
CC compound which modulates the biological activity of a receptor  
CC comprising (a) identifying peptides which bind the receptor by the  
CC method above, (b) using a number of the peptides to predict the receptor-  
CC modulating activity of a compound by (i) providing a panel of

CC identified peptides, where the members differ in their ability to bind  
CC to the receptor depending on reference conformations the receptor is  
CC in, where the effect of a number of reference substances known to  
CC modulate the biological activity of the receptor on the binding of each  
CC member of the panel is known and is characterised as a reference  
CC fingerprint for each reference substance, (ii) screening a test substance  
CC of unknown activity relative to the receptor to determine its effect on  
CC the binding of each member of the panel to the receptor, thereby  
CC obtaining a test fingerprint for the test substance, (iii) comparing the  
CC test fingerprint to the reference fingerprints and (iv) predicting the  
CC biological activity of the test substance based on the assumption that  
CC its biological activity will be similar to that of reference substances  
CC with similar fingerprints. The method is useful for identifying a binding  
CC peptide which binds a vertebrate, mammalian, preferably human receptor,  
CC an intracellular, nuclear, oestrogen or androgen receptor. The identified  
CC peptides which bind to the receptor are useful for predicting the  
CC receptor-modulating activity of a compound (e.g. ant/agonists).  
CC The receptor-binding library members are useful in the prediction of the  
CC ability of small organic molecules, suitable for pharmaceutical use  
CC (e.g. in the case of oestrogen receptors, for breast cancer treatment),  
CC to interact with the receptor. The analyte-binding molecules can also be  
CC used for in vivo imaging. The method has several advantages over whole  
CC animal-based assay systems in that the same technology can be applied to  
CC a variety of different receptors, the system can be used for high  
CC throughput screening and compound characterisation, and gives very  
CC distinct patterns for agonists and antagonists of receptor activity using  
CC very much less protein. The present sequence is an oestrogen receptor  
CC binding peptide from a combinatorial peptide library.  
XX  
SQ Sequence 15 AA;  
Query Match 83.9%; Score 26; DB 23; Length 15;  
Best Local Similarity 50.0%; Pred. No. 2.4e+02;  
Matches 3; Conservative 0; Mismatches 3; Indels 0; Gaps 0;  
QY 1 WXXWXP 6  
Db 8 WYDWTF 13  
RESULT 23  
AAR57391  
ID AAR57391 standard; Protein; 6 AA.  
XX  
AC AAR57391;  
XX  
DT 21-MAR-1995 (first entry)  
XX  
DE Peptide for treating diseases related to anti-DNA antibodies.  
XX  
KW Carrier; absorbing agent; treatment; anti-DNA antibody; immune complex.  
XX  
OS Synthetic.  
XX  
PN JP06192290-A.  
XX  
PD 12-JUL-1994.  
XX  
PF 18-JAN-1993; 93JP-0006098.  
XX  
PR 30-SEP-1992; 92JP-0261821.  
XX  
PA (KURS ) KURARAY CO LTD.  
XX  
DR WPI; 1994-260510/32.  
XX  
PT A peptide and an adsorbing agent prepd. by immobilising it on a  
PT carrier - useful for treatment of diseases related to anti-DNA  
PT antibodies and immune complexes  
XX  
PS Disclosure; Page 11; 14pp; Japanese.  
XX  
CC The sequences given in AAR57386-413 are peptides which are all covered



CC by the claimed generic formula:  
CC H-X-(A-B)n-Y-Z  
CC A = Trp, Phe or a peptide fragment consisting of 2 residues;  
CC B = Trp, Phe, Asn or Glu;  
CC X and Y = a bond or Asp, Glu, Arg, Lys, His or a peptide fragment  
CC consisting of 2-10 residues, provided that at least one of  
CC X or Y are present;  
CC Z = OH or NH2; and  
CC n = 2-5.  
CC These peptides may be immobilised on a carrier in the preparation of an  
CC absorbing agent which may be used in the treatment of diseases related  
CC to anti-DNA antibodies and/or immune complex.  
XX  
SQ Sequence 6 AA;  
  
Query Match 80.6%; Score 25; DB 15; Length 6;  
Best Local Similarity 50.0%; Pred. No. 9.3e+05;  
Matches 3; Conservative 0; Mismatches 3; Indels 0; Gaps 0;  
  
QY 1 WXXWXF 6  
| | | |  
Db 1 WFFWFF 6  
  
RESULT 24  
AAB01492  
ID AAB01492 standard; peptide; 6 AA.  
XX  
AC AAB01492;  
XX  
DT 08-NOV-2000 (first entry)  
XX  
DE Peptide which binds to transcription factor E2F-1 DNA binding domain.  
XX  
KW DNA binding; transcription factor; E2F; E2F-1; cell cycle; DP-1;  
KW activation; transcription; apoptosis; proliferative disorder;  
KW psoriasis; restenosis.  
XX  
OS Synthetic.  
XX  
PN WO200044771-A1.  
XX  
PD 03-AUG-2000.  
XX  
PF 26-JAN-2000; 2000WO-GB00227.  
XX  
PR 26-JAN-1999; 99GB-0001710.  
XX  
PA (PROL-) PROLIFIX LTD.  
XX  
PI Mueller R, Kontermann RE, Montigiani S;  
XX  
DR WPI; 2000-532806/48.  
XX  
PT Peptides binding to the DNA binding domain of transcription factor E2F  
PT and inhibiting cell cycle progression, useful for the treatment of  
PT cancer  
XX  
PS Claim 6; Page 2; 42pp; English.  
XX  
SQ Peptides which bind to the DNA binding domain of transcription  
CC factor E2F and inhibit cell cycle progression may be useful as  
CC research agents to investigate the interaction between E2F and DP-1,  
CC or the activation of transcription by E2F-1/DP-1 heterodimers. They  
CC may also be used for inducing apoptosis and/or cell cycle arrest in  
CC a cell, particularly for treatment of cancer or other proliferative  
CC disorders such as psoriasis and restenosis.  
XX  
SQ Sequence 6 AA;  
  
Query Match 80.6%; Score 25; DB 21; Length 6;  
Best Local Similarity 50.0%; Pred. No. 9.3e+05;  
Matches 3; Conservative 0; Mismatches 3; Indels 0; Gaps 0;

QY 1 WXXWXF 6  
| | | |  
Db 1 WVRWHF 6  
  
RESULT 25  
AAB01497  
ID AAB01497 standard; peptide; 6 AA.  
XX  
AC AAB01497;  
XX  
DT 08-NOV-2000 (first entry)  
XX  
DE Peptide which binds to transcription factor E2F-1 DNA binding domain.  
XX  
KW DNA binding; transcription factor; E2F; E2F-1; cell cycle; DP-1;  
KW activation; transcription; apoptosis; proliferative disorder;  
KW psoriasis; restenosis.  
XX  
OS Synthetic.  
XX  
PN WO200044771-A1.  
XX  
PD 03-AUG-2000.  
XX  
PF 26-JAN-2000; 2000WO-GB00227.  
XX  
PR 26-JAN-1999; 99GB-0001710.  
XX  
PA (PROL-) PROLIFIX LTD.  
XX  
PI Mueller R, Kontermann RE, Montigiani S;  
XX  
DR WPI; 2000-532806/48.  
XX  
PT Peptides binding to the DNA binding domain of transcription factor E2F  
PT and inhibiting cell cycle progression, useful for the treatment of  
PT cancer  
XX  
PS Claim 4; Page 9; 42pp; English.  
XX  
SQ Peptides which bind to the DNA binding domain of transcription  
CC factor E2F and inhibit cell cycle progression may be useful as  
CC research agents to investigate the interaction between E2F and DP-1,  
CC or the activation of transcription by E2F-1/DP-1 heterodimers. They  
CC may also be used for inducing apoptosis and/or cell cycle arrest in  
CC a cell, particularly for treatment of cancer or other proliferative  
CC disorders such as psoriasis and restenosis.  
XX  
SQ Sequence 6 AA;  
  
Query Match 80.6%; Score 25; DB 21; Length 6;  
Best Local Similarity 100.0%; Pred. No. 9.3e+05;  
Matches 6; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  
  
QY 1 WXXWXF 6  
| | | | |  
Db 1 WXXWXF 6  
  
RESULT 26  
AAB01499  
ID AAB01499 standard; peptide; 6 AA.  
XX

AC AAB01499;  
 XX  
 DT 08-NOV-2000 (first entry)  
 XX  
 DE Peptide which binds to transcription factor E2F-1 DNA binding domain.  
 XX  
 KW DNA binding; transcription factor; E2F; E2F-1; cell cycle; DP-1;  
 KW activation; transcription; apoptosis; proliferative disorder;  
 KW psoriasis; restenosis.  
 XX  
 OS Synthetic.  
 XX  
 FH Key Location/Qualifiers  
 FT Misc-difference 2  
 FT /note= "Any amino acid"  
 FT Misc-difference 3  
 FT /note= "Any amino acid"  
 XX  
 PN WO200044771-A1.  
 XX  
 PD 03-AUG-2000.  
 XX  
 PF 26-JAN-2000; 2000WO-GB00227.  
 XX  
 PR 26-JAN-1999; 99GB-0001710.  
 XX  
 PA (PROL-) PROLIFIX LTD.  
 XX  
 PI Mueller R, Kontermann RE, Montigiani S;  
 XX  
 DR WPI; 2000-532806/48.  
 XX  
 PT Peptides binding to the DNA binding domain of transcription factor E2F  
 PT and inhibiting cell cycle progression, useful for the treatment of  
 PT cancer  
 XX  
 PS Claim 4; Page 9; 42pp; English.  
 XX  
 CC Peptides which bind to the DNA binding domain of transcription  
 CC factor E2F and inhibit cell cycle progression may be useful as  
 CC research agents to investigate the interaction between E2F and DP-1,  
 CC or the activation of transcription by E2F-1/DP-1 heterodimers. They  
 CC may also be used for inducing apoptosis and/or cell cycle arrest in  
 CC a cell, particularly for treatment of cancer or other proliferative  
 CC disorders such as psoriasis and restenosis.  
 XX  
 SQ Sequence 6 AA;  
 Query Match 80.6%; Score 25; DB 21; Length 6;  
 Best Local Similarity 83.3%; Pred. No. 9.3e+05;  
 Matches 5; Conservative 0; Mismatches 1; Indels 0; Gaps 0;  
 QY 1 WXXWXXF 6  
 Db 1 WXXWTF 6  
 RESULT 27  
 ABR44865  
 ID ABR44865 standard; Peptide; 6 AA.  
 XX  
 AC ABR44865;  
 XX  
 DT 10-JUN-2003 (first entry)  
 XX  
 DE Staphylococcus aureus CHIPS-related peptide #55.  
 KW CHIPS; Chemotaxis Inhibitory Protein; C5a-receptor; C5aR;  
 KW formylated peptide receptor; FPR; neutrophil; monocyte; endothelial cell;  
 KW inflammation; cardiovascular disease; central nervous system disease;  
 KW gastrointestinal disease; skin disease; genitourinary disease;  
 KW joint disease; respiratory disease; HIV infection; antiinflammatory;  
 KW cardiant; cerebroprotective; neuroprotective; nootropic; dermatological;

KW gynecological; immunosuppressive; anti-HIV.  
 XX  
 OS Staphylococcus aureus.  
 OS Synthetic.  
 XX  
 PN WO2003006048-A1.  
 XX  
 PD 23-JAN-2003.  
 XX  
 PF 11-JUL-2001; 2001WO-EP08004.  
 XX  
 PR 11-JUL-2001; 2001WO-EP08004.  
 XX  
 PA (JARI-) JARI PHARM BV.  
 XX  
 PI Van Kessel CPM, Gosselaar-de Haas CJC, Kruijtzer JAW;  
 PI Van Strijp JAG;  
 XX  
 DR WPI; 2003-247783/25.  
 XX  
 PT Combination of peptides derived from chemotaxis inhibiting protein from  
 PT Staphylococcus aureus (CHIPS) having CHIPS activity, useful in  
 PT prophylaxis and treatment of inflammation, cardiovascular, skin and  
 PT kidney diseases  
 XX  
 PS Disclosure; Page 10; 89pp; English.  
 XX  
 CC The present invention relates to peptides (ABR44811-ABR47162 and  
 CC ABR47164-ABR47385) derived from the Chemotaxis Inhibitory Protein (CHIPS)  
 CC from Staphylococcus aureus. The peptide fragments are useful in the  
 CC prophylaxis or treatment of diseases or disorders involving the  
 CC C5a-receptor (C5aR) and/or formylated peptide receptor (FPR) or  
 CC neutrophils, monocytes and endothelial cells or involving acute or  
 CC chronic inflammation reactions. The diseases or disorders include  
 CC cardiovascular diseases, disease of the central nervous system,  
 CC gastrointestinal diseases, skin diseases, genitourinary diseases, joint  
 CC diseases, respiratory diseases and HIV infection.  
 XX  
 SQ Sequence 6 AA;  
 Query Match 80.6%; Score 25; DB 24; Length 6;  
 Best Local Similarity 50.0%; Pred. No. 9.3e+05;  
 Matches 3; Conservative 0; Mismatches 3; Indels 0; Gaps 0;  
 QY 1 WXXWXXF 6  
 Db 1 WSWFWPF 6  
 RESULT 28  
 ABR44866  
 ID ABR44866 standard; Peptide; 6 AA.  
 XX  
 AC ABR44866;  
 XX  
 DT 10-JUN-2003 (first entry)  
 XX  
 DE Staphylococcus aureus CHIPS-related peptide #56.  
 KW CHIPS; Chemotaxis Inhibitory Protein; C5a-receptor; C5aR;  
 KW formylated peptide receptor; FPR; neutrophil; monocyte; endothelial cell;  
 KW inflammation; cardiovascular disease; central nervous system disease;  
 KW gastrointestinal disease; skin disease; genitourinary disease;  
 KW joint disease; respiratory disease; HIV infection; antiinflammatory;  
 KW cardiant; cerebroprotective; neuroprotective; nootropic; dermatological;  
 KW gynecological; immunosuppressive; anti-HIV.  
 XX  
 OS Staphylococcus aureus.  
 OS Synthetic.  
 XX  
 PN WO2003006048-A1.  
 XX  
 PD 23-JAN-2003.

XX 11-JUL-2001; 2001WO-EP080004.  
PF 11-JUL-2001; 2001WO-EP080004.  
XX (JARI-) JARI PHARM BV.  
XX Van Kessel CPM, Gosselaar-de Haas CJC, Kruijtzer JAW;  
PI Van Strijp JAG;  
XX WPI; 2003-247783/25.  
DR  
XX Combination of peptides derived from chemotaxis inhibiting protein from  
PT Staphylococcus aureus (CHIPS) having CHIPS activity, useful in  
PT prophylaxis and treatment of inflammation, cardiovascular, skin and  
PT kidney diseases -  
XX  
PS Disclosure; Page 10; 89pp; English.  
XX The present invention relates to peptides (ABR44811-ABR47162 and  
CC ABR47164-ABR47385) derived from the Chemotaxis Inhibitory Protein (CHIPS)  
CC from Staphylococcus aureus. The peptide fragments are useful in the  
CC prophylaxis or treatment of diseases or disorders involving the  
CC C5a-receptor (C5aR) and/or formylated peptide receptor (FPR) or  
CC neutrophils, monocytes and endothelial cells or involving acute or  
CC chronic inflammation reactions. The diseases or disorders include  
CC cardiovascular diseases, disease of the central nervous system,  
CC gastrointestinal diseases, skin diseases, genitourinary diseases, joint  
CC diseases, respiratory diseases and HIV infection.  
XX  
SQ Sequence 6 AA;  
Query Match 80.6%; Score 25; DB 24; Length 6;  
Best Local Similarity 50.0%; Pred. No. 9.3e+05;  
Matches 3; Conservative 0; Mismatches 3; Indels 0; Gaps 0;  
QY 1 WXXWXF 6  
Db 1 WTFWPF 6  
RESULT 29  
ABR45311  
ID ABR45311 standard; Peptide; 6 AA.  
AC ABR45311;  
XX 10-JUN-2003 (first entry)  
DT  
XX Staphylococcus aureus CHIPS-related peptide #501.  
DE  
XX CHIPS; Chemotaxis Inhibitory Protein; C5a-receptor; C5aR;  
KW formylated peptide receptor; FPR; neutrophil; monocyte; endothelial cell;  
KW inflammation; cardiovascular disease; central nervous system disease;  
KW gastrointestinal disease; skin disease; genitourinary disease;  
KW joint disease; respiratory disease; HIV infection; antiinflammatory;  
KW cardiant; cerebroprotective; neuroprotective; nootropic; dermatological;  
KW gynecological; immunosuppressive; anti-HIV.  
XX  
OS Staphylococcus aureus.  
OS Synthetic.  
XX  
PN WO2003006048-A1.  
XX 23-JAN-2003.  
PD  
XX 11-JUL-2001; 2001WO-EP080004.  
PF  
XX 11-JUL-2001; 2001WO-EP080004.  
PR (JARI-) JARI PHARM BV.  
XX Van Kessel CPM, Gosselaar-de Haas CJC, Kruijtzer JAW;  
PI

PI Van Strijp JAG;  
XX WPI; 2003-247783/25.  
DR  
XX Combination of peptides derived from chemotaxis inhibiting protein from  
PT Staphylococcus aureus (CHIPS) having CHIPS activity, useful in  
PT prophylaxis and treatment of inflammation, cardiovascular, skin and  
PT kidney diseases -  
XX  
PS Disclosure; Page 12; 89pp; English.  
XX The present invention relates to peptides (ABR44811-ABR47162 and  
CC ABR47164-ABR47385) derived from the Chemotaxis Inhibitory Protein (CHIPS)  
CC from Staphylococcus aureus. The peptide fragments are useful in the  
CC prophylaxis or treatment of diseases or disorders involving the  
CC C5a-receptor (C5aR) and/or formylated peptide receptor (FPR) or  
CC neutrophils, monocytes and endothelial cells or involving acute or  
CC chronic inflammation reactions. The diseases or disorders include  
CC cardiovascular diseases, disease of the central nervous system,  
CC gastrointestinal diseases, skin diseases, genitourinary diseases, joint  
CC diseases, respiratory diseases and HIV infection.  
XX  
SQ Sequence 6 AA;  
Query Match 80.6%; Score 25; DB 24; Length 6;  
Best Local Similarity 50.0%; Pred. No. 9.3e+05;  
Matches 3; Conservative 0; Mismatches 3; Indels 0; Gaps 0;  
QY 1 WXXWXF 6  
Db 1 WFFWFF 6  
RESULT 30  
ABR45312  
ID ABR45312 standard; Peptide; 6 AA.  
XX  
AC ABR45312;  
XX 10-JUN-2003 (first entry)  
DT  
XX Staphylococcus aureus CHIPS-related peptide #502.  
DE  
XX CHIPS; Chemotaxis Inhibitory Protein; C5a-receptor; C5aR;  
KW formylated peptide receptor; FPR; neutrophil; monocyte; endothelial cell;  
KW inflammation; cardiovascular disease; central nervous system disease;  
KW gastrointestinal disease; skin disease; genitourinary disease;  
KW joint disease; respiratory disease; HIV infection; antiinflammatory;  
KW cardiant; cerebroprotective; neuroprotective; nootropic; dermatological;  
KW gynecological; immunosuppressive; anti-HIV.  
XX  
OS Staphylococcus aureus.  
OS Synthetic.  
XX  
PN WO2003006048-A1.  
XX 23-JAN-2003.  
PD  
XX 11-JUL-2001; 2001WO-EP080004.  
PF  
XX 11-JUL-2001; 2001WO-EP080004.  
PR (JARI-) JARI PHARM BV.  
XX  
PI Van Kessel CPM, Gosselaar-de Haas CJC, Kruijtzer JAW;  
PI Van Strijp JAG;  
XX WPI; 2003-247783/25.  
DR  
XX Combination of peptides derived from chemotaxis inhibiting protein from  
PT Staphylococcus aureus (CHIPS) having CHIPS activity, useful in  
PT prophylaxis and treatment of inflammation, cardiovascular, skin and  
PT kidney diseases -  
PT

XX Disclosure; Page 12; 89pp; English.

PS

XX The present invention relates to peptides (ABR44811-ABR47162 and ABR47164-ABR47385) derived from the Chemotaxis Inhibitory Protein (CHIPS) from Staphylococcus aureus. The peptide fragments are useful in the prophylaxis or treatment of diseases or disorders involving the C5a-receptor (C5aR) and/or formylated peptide receptor (FPR) or neutrophils, monocytes and endothelial cells or involving acute or chronic inflammation reactions. The diseases or disorders include cardiovascular diseases, disease of the central nervous system, gastrointestinal diseases, skin diseases, genitourinary diseases, joint diseases, respiratory diseases and HIV infection.

XX

SQ Sequence 6 AA;

Query Match 80.6%; Score 25; DB 24; Length 6;  
Best Local Similarity 50.0%; Pred. No. 9.3e+05;  
Matches 3; Conservative 0; Mismatches 3; Indels 0; Gaps 0;

QY 1 WXXWXF 6  
| | |  
Db 1 WIFWFF 6

RESULT 31

ABR45367

ID ABR45367 standard; Peptide; 6 AA.

XX

AC ABR45367;

XX

DT 10-JUN-2003 (first entry)

XX

DE Staphylococcus aureus CHIPS-related peptide #557.

XX

KW CHIPS; Chemotaxis Inhibitory Protein; C5a-receptor; C5aR; formylated peptide receptor; FPR; neutrophil; monocyte; endothelial cell; inflammation; cardiovascular disease; central nervous system disease; gastrointestinal disease; skin disease; genitourinary disease; joint disease; respiratory disease; HIV infection; antiinflammatory; cardiant; cerebroprotective; neuroprotective; nootropic; dermatological; gynecological; immunosuppressive; anti-HIV.

XX

OS Staphylococcus aureus.

OS Synthetic.

XX WO2003006048-A1.

XX

PN 23-JAN-2003.

XX

PF 11-JUL-2001; 2001WO-EP08004.

XX

PR 11-JUL-2001; 2001WO-EP08004.

XX

PA (JARI-) JARI PHARM BV.

XX

PI Van Kessel CPM, Gosselaar-de Haas CJC, Kruijtzer JAW;  
PI Van Strijp JAG;

XX

DR WPI; 2003-247783/25.

XX

PT Combination of peptides derived from chemotaxis inhibiting protein from Staphylococcus aureus (CHIPS) having CHIPS activity, useful in prophylaxis and treatment of inflammation, cardiovascular, skin and kidney diseases -

XX

PS Disclosure; Page 12; 89pp; English.

XX

CC The present invention relates to peptides (ABR44811-ABR47162 and ABR47164-ABR47385) derived from the Chemotaxis Inhibitory Protein (CHIPS) from Staphylococcus aureus. The peptide fragments are useful in the prophylaxis or treatment of diseases or disorders involving the C5a-receptor (C5aR) and/or formylated peptide receptor (FPR) or neutrophils, monocytes and endothelial cells or involving acute or chronic inflammation reactions. The diseases or disorders include cardiovascular diseases, disease of the central nervous system, gastrointestinal diseases, skin diseases, genitourinary diseases, joint diseases, respiratory diseases and HIV infection.

XX

CC

CC neutrophils, monocytes and endothelial cells or involving acute or chronic inflammation reactions. The diseases or disorders include cardiovascular diseases, disease of the central nervous system, gastrointestinal diseases, skin diseases, genitourinary diseases, joint diseases, respiratory diseases and HIV infection.

XX

SQ Sequence 6 AA;

Query Match 80.6%; Score 25; DB 24; Length 6;  
Best Local Similarity 50.0%; Pred. No. 9.3e+05;  
Matches 3; Conservative 0; Mismatches 3; Indels 0; Gaps 0;

QY 1 WXXWXF 6  
| | |  
Db 1 WFFWIF 6

RESULT 32

ABR45368

ID ABR45368 standard; Peptide; 6 AA.

XX

AC ABR45368;

XX

DT 10-JUN-2003 (first entry)

XX

DE Staphylococcus aureus CHIPS-related peptide #558.

XX

KW CHIPS; Chemotaxis Inhibitory Protein; C5a-receptor; C5aR; formylated peptide receptor; FPR; neutrophil; monocyte; endothelial cell; inflammation; cardiovascular disease; central nervous system disease; gastrointestinal disease; skin disease; genitourinary disease; joint disease; respiratory disease; HIV infection; antiinflammatory; cardiant; cerebroprotective; neuroprotective; nootropic; dermatological; gynecological; immunosuppressive; anti-HIV.

XX

OS Staphylococcus aureus.

OS Synthetic.

XX WO2003006048-A1.

XX

PN 23-JAN-2003.

XX

PF 11-JUL-2001; 2001WO-EP08004.

XX

PR 11-JUL-2001; 2001WO-EP08004.

XX

PA (JARI-) JARI PHARM BV.

XX

PI Van Kessel CPM, Gosselaar-de Haas CJC, Kruijtzer JAW;  
PI Van Strijp JAG;

XX

DR WPI; 2003-247783/25.

XX

PT Combination of peptides derived from chemotaxis inhibiting protein from Staphylococcus aureus (CHIPS) having CHIPS activity, useful in prophylaxis and treatment of inflammation, cardiovascular, skin and kidney diseases -

XX

PS Disclosure; Page 12; 89pp; English.

XX

CC The present invention relates to peptides (ABR44811-ABR47162 and ABR47164-ABR47385) derived from the Chemotaxis Inhibitory Protein (CHIPS) from Staphylococcus aureus. The peptide fragments are useful in the prophylaxis or treatment of diseases or disorders involving the C5a-receptor (C5aR) and/or formylated peptide receptor (FPR) or neutrophils, monocytes and endothelial cells or involving acute or chronic inflammation reactions. The diseases or disorders include cardiovascular diseases, disease of the central nervous system, gastrointestinal diseases, skin diseases, genitourinary diseases, joint diseases, respiratory diseases and HIV infection.

XX

CC

Sequence 6 AA;



Query Match 80.6%; Score 25; DB 24; Length 6;  
Best Local Similarity 50.0%; Pred. No. 9.3e+05;  
Matches 3; Conservative 0; Mismatches 3; Indels 0; Gaps 0;

QY 1 WXXWXF 6  
| | | |  
Db 1 WIFWIF 6

RESULT 33  
ABR45423  
ID ABR45423 standard; Peptide; 6 AA.  
XX  
AC ABR45423;  
XX  
DT 10-JUN-2003 (first entry)  
XX  
DE Staphylococcus aureus CHIPS-related peptide #613.  
XX  
KW CHIPS; Chemotaxis Inhibitory Protein; C5a-receptor; C5aR;  
KW formylated peptide receptor; FPR; neutrophil; monocyte; endothelial cell;  
KW inflammation; cardiovascular disease; central nervous system disease;  
KW gastrointestinal disease; skin disease; genitourinary disease;  
KW joint disease; respiratory disease; HIV infection; antiinflammatory;  
KW cardiant; cerebroprotective; neuroprotective; nootropic; dermatological;  
KW gynecological; immunosuppressive; anti-HIV.  
XX  
OS Staphylococcus aureus.  
OS Synthetic.  
XX  
PN WO2003006048-A1.  
XX  
PD 23-JAN-2003.  
XX  
PF 11-JUL-2001; 2001WO-EP08004.  
XX  
PR 11-JUL-2001; 2001WO-EP08004.  
XX  
PA (JARI-) JARI PHARM BV.  
XX  
PI Van Kessel CPM, Gosselaar-de Haas CJC, Kruijtzer JAW;  
PI Van Strijp JAG;  
XX  
DR WPI; 2003-247783/25.  
XX  
PT Combination of peptides derived from chemotaxis inhibiting protein from  
PT Staphylococcus aureus (CHIPS) having CHIPS activity, useful in  
PT prophylaxis and treatment of inflammation, cardiovascular, skin and  
PT kidney diseases -  
XX  
PS Disclosure; Page 12; 89pp; English.  
XX  
CC The present invention relates to peptides (ABR44811-ABR47162 and  
CC ABR47164-ABR47385) derived from the Chemotaxis Inhibitory Protein (CHIPS)  
CC from Staphylococcus aureus. The peptide fragments are useful in the  
CC prophylaxis or treatment of diseases or disorders involving the  
CC C5a-receptor (C5aR) and/or formylated peptide receptor (FPR) or  
CC neutrophils, monocytes and endothelial cells or involving acute or  
CC chronic inflammation reactions. The diseases or disorders include  
CC cardiovascular diseases, disease of the central nervous system,  
CC gastrointestinal diseases, skin diseases, genitourinary diseases, joint  
CC diseases, respiratory diseases and HIV infection.

Sequence 6 AA;  
Query Match 80.6%; Score 25; DB 24; Length 6;  
Best Local Similarity 50.0%; Pred. No. 9.3e+05;  
Matches 3; Conservative 0; Mismatches 3; Indels 0; Gaps 0;

QY 1 WXXWXF 6  
| | | |  
Db 1 WFFWLF 6

RESULT 34  
ABR45424  
ID ABR45424 standard; Peptide; 6 AA.  
XX  
AC ABR45424;  
XX  
DT 10-JUN-2003 (first entry)  
XX  
DE Staphylococcus aureus CHIPS-related peptide #614.  
XX  
KW CHIPS; Chemotaxis Inhibitory Protein; C5a-receptor; C5aR;  
KW formylated peptide receptor; FPR; neutrophil; monocyte; endothelial cell;  
KW inflammation; cardiovascular disease; central nervous system disease;  
KW gastrointestinal disease; skin disease; genitourinary disease;  
KW joint disease; respiratory disease; HIV infection; antiinflammatory;  
KW cardiant; cerebroprotective; neuroprotective; nootropic; dermatological;  
KW gynecological; immunosuppressive; anti-HIV.  
XX  
OS Staphylococcus aureus.  
OS Synthetic.  
XX  
PN WO2003006048-A1.  
XX  
PD 23-JAN-2003.  
XX  
PF 11-JUL-2001; 2001WO-EP08004.  
XX  
PR 11-JUL-2001; 2001WO-EP08004.  
XX  
PA (JARI-) JARI PHARM BV.  
XX  
PI Van Kessel CPM, Gosselaar-de Haas CJC, Kruijtzer JAW;  
PI Van Strijp JAG;  
XX  
DR WPI; 2003-247783/25.  
XX  
PT Combination of peptides derived from chemotaxis inhibiting protein from  
PT Staphylococcus aureus (CHIPS) having CHIPS activity, useful in  
PT prophylaxis and treatment of inflammation, cardiovascular, skin and  
PT kidney diseases -  
XX  
PS Disclosure; Page 12; 89pp; English.  
XX  
CC The present invention relates to peptides (ABR44811-ABR47162 and  
CC ABR47164-ABR47385) derived from the Chemotaxis Inhibitory Protein (CHIPS)  
CC from Staphylococcus aureus. The peptide fragments are useful in the  
CC prophylaxis or treatment of diseases or disorders involving the  
CC C5a-receptor (C5aR) and/or formylated peptide receptor (FPR) or  
CC neutrophils, monocytes and endothelial cells or involving acute or  
CC chronic inflammation reactions. The diseases or disorders include  
CC cardiovascular diseases, disease of the central nervous system,  
CC gastrointestinal diseases, skin diseases, genitourinary diseases, joint  
CC diseases, respiratory diseases and HIV infection.

Sequence 6 AA;  
Query Match 80.6%; Score 25; DB 24; Length 6;  
Best Local Similarity 50.0%; Pred. No. 9.3e+05;  
Matches 3; Conservative 0; Mismatches 3; Indels 0; Gaps 0;

QY 1 WXXWXF 6  
| | | |  
Db 1 WIFWLF 6

RESULT 35  
ABR45479  
ID ABR45479 standard; Peptide; 6 AA.  
XX  
AC ABR45479;  
XX  
DT 10-JUN-2003 (first entry)



XX Staphylococcus aureus CHIPS-related peptide #669.

DE CHIPS; Chemotaxis Inhibitory Protein; C5a-receptor; C5aR;

XX formylated peptide receptor; FPR; neutrophil; monocyte; endothelial cell;

KW inflammation; cardiovascular disease; central nervous system disease;

KW gastrointestinal disease; skin disease; genitourinary disease;

KW joint disease; respiratory disease; HIV infection; antiinflammatory;

KW cardiant; cerebroprotective; neuroprotective; nootropic; dermatological;

KW gynecological; immunosuppressive; anti-HIV.

XX Staphylococcus aureus.

OS Staphylococcus aureus.

OS Synthetic.

XX WO2003006048-A1.

PN 23-JAN-2003.

XX 11-JUL-2001; 2001WO-EP08004.

XX 11-JUL-2001; 2001WO-EP08004.

PR (JARI-) JARI PHARM BV.

XX Van Kessel CPM, Gosselaar-de Haas CJC, Kruijtzer JAW;

PI Van Strijp JAG;

PI WPI; 2003-247783/25.

DR Combination of peptides derived from chemotaxis inhibiting protein from

XX Staphylococcus aureus (CHIPS) having CHIPS activity, useful in

PT prophylaxis and treatment of inflammation, cardiovascular, skin and

PT kidney diseases -

PT Disclosure; Page 13; 89pp; English.

PS The present invention relates to peptides (ABR44811-ABR47162 and

XX ABR47164-ABR47385) derived from the Chemotaxis Inhibitory Protein (CHIPS)

CC from Staphylococcus aureus. The peptide fragments are useful in the

CC prophylaxis or treatment of diseases or disorders involving the

CC C5a-receptor (C5aR) and/or formylated peptide receptor (FPR) or

CC neutrophils, monocytes and endothelial cells or involving acute or

CC chronic inflammation reactions. The diseases or disorders include

CC cardiovascular diseases, disease of the central nervous system,

CC gastrointestinal diseases, skin diseases, genitourinary diseases, joint

CC diseases, respiratory diseases and HIV infection.

XX Sequence 6 AA;

SQ Query Match 80.6%; Score 25; DB 24; Length 6;

Best Local Similarity 50.0%; Pred. No. 9.3e+05;

Matches 3; Conservative 0; Mismatches 3; Indels 0; Gaps 0;

QY 1 WXXWXF 6

Db 1 WFFWVF 6

RESULT 36

ABR45480

ID ABR45480 standard; Peptide; 6 AA.

XX ABR45480;

AC 10-JUN-2003 (first entry)

XX Staphylococcus aureus CHIPS-related peptide #670.

DE CHIPS; Chemotaxis Inhibitory Protein; C5a-receptor; C5aR;

XX formylated peptide receptor; FPR; neutrophil; monocyte; endothelial cell;

KW inflammation; cardiovascular disease; central nervous system disease;

KW gastrointestinal disease; skin disease; genitourinary disease;

KW joint disease; respiratory disease; HIV infection; antiinflammatory;

KW cardiant; cerebroprotective; neuroprotective; nootropic; dermatological;

KW gynecological; immunosuppressive; anti-HIV.

XX Staphylococcus aureus.

OS Synthetic.

XX WO2003006048-A1.

PN 23-JAN-2003.

XX 11-JUL-2001; 2001WO-EP08004.

XX 11-JUL-2001; 2001WO-EP08004.

PR (JARI-) JARI PHARM BV.

XX Van Kessel CPM, Gosselaar-de Haas CJC, Kruijtzer JAW;

PI Van Strijp JAG;

PI WPI; 2003-247783/25.

DR Combination of peptides derived from chemotaxis inhibiting protein from

XX Staphylococcus aureus (CHIPS) having CHIPS activity, useful in

PT prophylaxis and treatment of inflammation, cardiovascular, skin and

PT kidney diseases -

PT Disclosure; Page 13; 89pp; English.

PS The present invention relates to peptides (ABR44811-ABR47162 and

XX ABR47164-ABR47385) derived from the Chemotaxis Inhibitory Protein (CHIPS)

CC from Staphylococcus aureus. The peptide fragments are useful in the

CC prophylaxis or treatment of diseases or disorders involving the

CC C5a-receptor (C5aR) and/or formylated peptide receptor (FPR) or

CC neutrophils, monocytes and endothelial cells or involving acute or

CC chronic inflammation reactions. The diseases or disorders include

CC cardiovascular diseases, disease of the central nervous system,

CC gastrointestinal diseases, skin diseases, genitourinary diseases, joint

CC diseases, respiratory diseases and HIV infection.

XX Sequence 6 AA;

SQ Query Match 80.6%; Score 25; DB 24; Length 6;

Best Local Similarity 50.0%; Pred. No. 9.3e+05;

Matches 3; Conservative 0; Mismatches 3; Indels 0; Gaps 0;

QY 1 WXXWXF 6

Db 1 WFFWVF 6

RESULT 37

ABR45537

ID ABR45537 standard; Peptide; 6 AA.

XX ABR45537;

AC 10-JUN-2003 (first entry)

XX Staphylococcus aureus CHIPS-related peptide #727.

DE CHIPS; Chemotaxis Inhibitory Protein; C5a-receptor; C5aR;

XX formylated peptide receptor; FPR; neutrophil; monocyte; endothelial cell;

KW inflammation; cardiovascular disease; central nervous system disease;

KW gastrointestinal disease; skin disease; genitourinary disease;

KW joint disease; respiratory disease; HIV infection; antiinflammatory;

KW cardiant; cerebroprotective; neuroprotective; nootropic; dermatological;

KW gynecological; immunosuppressive; anti-HIV.

XX Staphylococcus aureus.

OS Synthetic.

XX WO2003006048-A1.

PD 23-JAN-2003.  
XX  
PF 11-JUL-2001; 2001WO-EP08004.  
XX  
PR 11-JUL-2001; 2001WO-EP08004.  
XX  
PA (JARI-) JARI PHARM BV.  
XX  
PI Van Kessel CPM, Gosselaar-de Haas CJC, Kruijtzer JAW;  
PI Van Strijp JAG;  
XX  
DR WPI; 2003-247783/25.  
XX  
XX Combination of peptides derived from chemotaxis inhibiting protein from  
PT Staphylococcus aureus (CHIPS) having CHIPS activity, useful in  
PT prophylaxis and treatment of inflammation, cardiovascular, skin and  
PT kidney diseases -  
XX  
XX Disclosure; Page 13; 89pp; English.  
PS  
XX  
XX The present invention relates to peptides (ABR44811-ABR47162 and  
CC ABR47164-ABR47385) derived from the Chemotaxis Inhibitory Protein (CHIPS)  
CC from Staphylococcus aureus. The peptide fragments are useful in the  
CC prophylaxis or treatment of diseases or disorders involving the  
CC C5a-receptor (C5aR) and/or formylated peptide receptor (FPR) or  
CC neutrophils, monocytes and endothelial cells or involving acute or  
CC chronic inflammation reactions. The diseases or disorders include  
CC cardiovascular diseases, disease of the central nervous system,  
CC gastrointestinal diseases, skin diseases, genitourinary diseases, joint  
CC diseases, respiratory diseases and HIV infection.  
XX  
SQ Sequence 6 AA;  
  
Query Match 80.6%; Score 25; DB 24; Length 6;  
Best Local Similarity 50.0%; Pred. No. 9.3e+05;  
Matches 3; Conservative 0; Mismatches 3; Indels 0; Gaps 0;  
  
QY 1 WXXWXF 6  
| | |  
Db 1 WSFWWF 6  
  
RESULT 38  
ABR45538  
ID ABR45538 standard; Peptide; 6 AA.  
XX  
AC ABR45538;  
XX  
DT 10-JUN-2003 (first entry)  
XX  
DE Staphylococcus aureus CHIPS-related peptide #728.  
XX  
KW CHIPS; Chemotaxis Inhibitory Protein; C5a-receptor; C5aR;  
KW formylated peptide receptor; FPR; neutrophil; monocyte; endothelial cell;  
KW inflammation; cardiovascular disease; central nervous system disease;  
KW gastrointestinal disease; skin disease; genitourinary disease;  
KW joint disease; respiratory disease; HIV infection; antiinflammatory;  
KW cardiant; cerebroprotective; neuroprotective; nootropic; dermatological;  
KW gynecological; immunosuppressive; anti-HIV.  
XX  
OS Staphylococcus aureus.  
OS Synthetic.  
XX  
PN WO2003006048-A1.  
XX  
PD 23-JAN-2003.  
XX  
PF 11-JUL-2001; 2001WO-EP08004.  
XX  
PR 11-JUL-2001; 2001WO-EP08004.  
XX  
PA (JARI-) JARI PHARM BV.  
XX

PI Van Kessel CPM, Gosselaar-de Haas CJC, Kruijtzer JAW;  
PI Van Strijp JAG;  
XX  
DR WPI; 2003-247783/25.  
XX  
XX Combination of peptides derived from chemotaxis inhibiting protein from  
PT Staphylococcus aureus (CHIPS) having CHIPS activity, useful in  
PT prophylaxis and treatment of inflammation, cardiovascular, skin and  
PT kidney diseases -  
XX  
XX Disclosure; Page 13; 89pp; English.  
PS  
XX  
XX The present invention relates to peptides (ABR44811-ABR47162 and  
CC ABR47164-ABR47385) derived from the Chemotaxis Inhibitory Protein (CHIPS)  
CC from Staphylococcus aureus. The peptide fragments are useful in the  
CC prophylaxis or treatment of diseases or disorders involving the  
CC C5a-receptor (C5aR) and/or formylated peptide receptor (FPR) or  
CC neutrophils, monocytes and endothelial cells or involving acute or  
CC chronic inflammation reactions. The diseases or disorders include  
CC cardiovascular diseases, disease of the central nervous system,  
CC gastrointestinal diseases, skin diseases, genitourinary diseases, joint  
CC diseases, respiratory diseases and HIV infection.  
XX  
SQ Sequence 6 AA;  
  
Query Match 80.6%; Score 25; DB 24; Length 6;  
Best Local Similarity 50.0%; Pred. No. 9.3e+05;  
Matches 3; Conservative 0; Mismatches 3; Indels 0; Gaps 0;  
  
QY 1 WXXWXF 6  
| | |  
Db 1 WTFWWF 6  
  
RESULT 39  
ABR45591  
ID ABR45591 standard; Peptide; 6 AA.  
XX  
AC ABR45591;  
XX  
DT 10-JUN-2003 (first entry)  
XX  
DE Staphylococcus aureus CHIPS-related peptide #781.  
XX  
KW CHIPS; Chemotaxis Inhibitory Protein; C5a-receptor; C5aR;  
KW formylated peptide receptor; FPR; neutrophil; monocyte; endothelial cell;  
KW inflammation; cardiovascular disease; central nervous system disease;  
KW gastrointestinal disease; skin disease; genitourinary disease;  
KW joint disease; respiratory disease; HIV infection; antiinflammatory;  
KW cardiant; cerebroprotective; neuroprotective; nootropic; dermatological;  
KW gynecological; immunosuppressive; anti-HIV.  
XX  
OS Staphylococcus aureus.  
OS Synthetic.  
XX  
PN WO2003006048-A1.  
XX  
PD 23-JAN-2003.  
XX  
PF 11-JUL-2001; 2001WO-EP08004.  
XX  
PR 11-JUL-2001; 2001WO-EP08004.  
XX  
PA (JARI-) JARI PHARM BV.  
XX  
PI Van Kessel CPM, Gosselaar-de Haas CJC, Kruijtzer JAW;  
PI Van Strijp JAG;  
XX  
DR WPI; 2003-247783/25.  
XX  
XX Combination of peptides derived from chemotaxis inhibiting protein from  
PT Staphylococcus aureus (CHIPS) having CHIPS activity, useful in  
PT prophylaxis and treatment of inflammation, cardiovascular, skin and  
PT

PT kidney diseases -  
PS Disclosure; Page 13; 89pp; English.  
XX  
CC The present invention relates to peptides (ABR44811-ABR47162 and  
CC ABR47164-ABR47385) derived from the Chemotaxis Inhibitory Protein (CHIPS)  
CC from Staphylococcus aureus. The peptide fragments are useful in the  
CC prophylaxis or treatment of diseases or disorders involving the  
CC C5a-receptor (C5aR) and/or formylated peptide receptor (FPR) or  
CC neutrophils, monocytes and endothelial cells or involving acute or  
CC chronic inflammation reactions. The diseases or disorders include  
CC cardiovascular diseases, disease of the central nervous system,  
CC gastrointestinal diseases, skin diseases, genitourinary diseases, joint  
CC diseases, respiratory diseases and HIV infection.  
XX  
SQ Sequence 6 AA;  
Query Match 80.6%; Score 25; DB 24; Length 6;  
Best Local Similarity 50.0%; Pred. No. 9.3e+05;  
Matches 3; Conservative 0; Mismatches 3; Indels 0; Gaps 0;  
QY 1 WXXWXF 6  
Db 1 WFFWYF 6  
RESULT 40  
ABR45592  
ID ABR45592 standard; Peptide; 6 AA.  
XX  
AC ABR45592;  
XX  
DT 10-JUN-2003 (first entry)  
XX  
DE Staphylococcus aureus CHIPS-related peptide #782.  
XX  
KW CHIPS; Chemotaxis Inhibitory Protein; C5a-receptor; C5aR;  
KW formylated peptide receptor; FPR; neutrophil; monocyte; endothelial cell;  
KW inflammation; cardiovascular disease; central nervous system disease;  
KW gastrointestinal disease; skin disease; genitourinary disease;  
KW joint disease; respiratory disease; HIV infection; antiinflammatory;  
KW cardiant; cerebroprotective; neuroprotective; nootropic; dermatological;  
KW gynecological; immunosuppressive; anti-HIV.  
XX  
OS Staphylococcus aureus.  
OS Synthetic.  
XX  
PN WO2003006048-A1.  
XX  
PD 23-JAN-2003.  
XX  
PF 11-JUL-2001; 2001WO-EP08004.  
XX  
PR 11-JUL-2001; 2001WO-EP08004.  
XX  
PA (JARI-) JARI PHARM BV.  
XX  
PI Van Kessel CPM, Gosselaar-de Haas CJC, Kruijtzer JAW;  
PI Van Strijp JAG;  
XX  
DR WPI; 2003-247783/25.  
XX  
PT Combination of peptides derived from chemotaxis inhibiting protein from  
PT Staphylococcus aureus (CHIPS) having CHIPS activity, useful in  
PT prophylaxis and treatment of inflammation, cardiovascular, skin and  
PT kidney diseases -  
XX  
PS Disclosure; Page 13; 89pp; English.  
XX  
CC The present invention relates to peptides (ABR44811-ABR47162 and  
CC ABR47164-ABR47385) derived from the Chemotaxis Inhibitory Protein (CHIPS)  
CC from Staphylococcus aureus. The peptide fragments are useful in the  
CC prophylaxis or treatment of diseases or disorders involving the

CC C5a-receptor (C5aR) and/or formylated peptide receptor (FPR) or  
CC neutrophils, monocytes and endothelial cells or involving acute or  
CC chronic inflammation reactions. The diseases or disorders include  
CC cardiovascular diseases, disease of the central nervous system,  
CC gastrointestinal diseases, skin diseases, genitourinary diseases, joint  
CC diseases, respiratory diseases and HIV infection.  
XX  
SQ Sequence 6 AA;  
Query Match 80.6%; Score 25; DB 24; Length 6;  
Best Local Similarity 50.0%; Pred. No. 9.3e+05;  
Matches 3; Conservative 0; Mismatches 3; Indels 0; Gaps 0;  
QY 1 WXXWXF 6  
Db 1 WFFWYF 6  
RESULT 41  
AAM45777  
ID AAM45777 standard; Peptide; 7 AA.  
XX  
AC AAM45777;  
XX  
DT 25-OCT-2001 (first entry)  
XX  
DE H11 binding site consensus conforming peptide (CCP) #2048.  
XX  
KW Antigen-binding; tumour; diagnosis; stress protein-peptide complex; SPPC;  
KW immunogenically cross-reactive; cancer; immunogenic cancer cell;  
KW cytostatic; vaccine; tumour-specific immunogenic response inducer;  
KW astrocytoma; fibrosarcoma; myxosarcoma; liposarcoma; oligodendroglioma;  
KW ependymoma; medulloblastoma; primitive neural ectodermal tumour.  
XX  
OS Homo sapiens.  
OS Synthetic.  
XX  
PN CA2290722-A1.  
XX  
PD 08-JUN-2001.  
XX  
PF 08-DEC-1999; 99CA-2290722.  
XX  
PR 08-DEC-1999; 99CA-2290722.  
XX  
PA (NOVO-) NOVOPHARM BIOTECH INC.  
XX  
PI Kaplan HA, Maiti PK, Fast DG, Herman W, Dan MD, Lewis KE;  
PI Entwistle JM, MacDonald GC;  
XX  
DR WPI; 2001-425937/46.  
XX  
PT Composition useful for treating and diagnosing cancer, comprises stress  
PT protein-peptide complexes associated with tumor, and isolated  
PT antigen-binding fragments of an antibody that binds specifically to the  
PT complex -  
XX  
PS Example 4; Page 108; 154pp; English.  
XX  
CC The present invention describes a composition (I) comprising stress  
CC protein-peptide complexes (SPPC) associated with tumours that is  
CC specifically immunogenically cross-reactive with cell surface-associated  
CC SPPCs specific to target cancer (TC). Also described is an isolated  
CC antigen-binding fragment of an antibody that binds specifically to SPPCs  
CC or a population of different SPPCs consisting of immunogenic cancer cell  
CC surface-associated SPPC of TC. (I) has cytostatic activity and can be  
CC used in vaccine production and as a tumour-specific immunogenic response  
CC inducer. (I) is useful for treating 71 types of cancers or tumours in a  
CC subject, such as astrocytoma, fibrosarcoma, myxosarcoma, liposarcoma,  
CC oligodendroglioma, ependymoma, medulloblastoma, and primitive neural  
CC ectodermal tumour (PNET). (I) is useful as cancer immunogen including  
CC vaccines. (I) is useful for diagnostic and palliative use, for detecting  
CC or imaging cancer cells, and to monitor the course of amelioration of

CC malignancy in an individual. AAM43707 to AAM47109 represent peptides  
CC which are used in the exemplification of the present invention.

XX  
SQ Sequence 7 AA;  
Query Match 80.6%; Score 25; DB 22; Length 7;  
Best Local Similarity 50.0%; Pred. No. 9.3e+05;  
Matches 3; Conservative 0; Mismatches 3; Indels 0; Gaps 0;  
QY 1 WXXWXF 6  
| | | |  
Db 1 WRRWNF 6

RESULT 42  
AAB01498  
ID AAB01498 standard; peptide; 9 AA.  
XX  
AC AAB01498;  
XX  
DT 08-NOV-2000 (first entry)  
XX  
DE Peptide which binds to transcription factor E2F-1 DNA binding domain.  
XX  
KW DNA binding; transcription factor; E2F; E2F-1; cell cycle; DP-1;  
KW activation; transcription; apoptosis; proliferative disorder;  
KW psoriasis; restenosis.  
XX  
OS Synthetic.

XX  
FH Key Location/Qualifiers  
FT Misc-difference 2 /note= "Any amino acid"  
FT Misc-difference 3 /note= "Any amino acid"  
FT Misc-difference 5 /note= "Any amino acid"  
FT Misc-difference 7 /note= "Any amino acid"  
FT Misc-difference 8 /note= "Any amino acid"  
FT Misc-difference /note= "Any amino acid"

XX  
PN WO200044771-A1.  
XX  
PD 03-AUG-2000.  
XX  
PF 26-JAN-2000; 2000WO-GB00227.  
XX  
PR 26-JAN-1999; 99GB-0001710.  
XX  
PA (PROL-) PROLIFIX LTD.  
XX  
PI Mueller R, Kontermann RE, Montigiani S;  
XX  
DR WPI; 2000-532806/48.  
XX  
PT Peptides binding to the DNA binding domain of transcription factor E2F  
PT and inhibiting cell cycle progression, useful for the treatment of  
PT cancer  
XX  
PS Claim 4; Page 9; 42pp; English.  
XX  
CC Peptides which bind to the DNA binding domain of transcription  
CC factor E2F and inhibit cell cycle progression may be useful as  
CC research agents to investigate the interaction between E2F and DP-1,  
CC or the activation of transcription by E2F-1/DP-1 heterodimers. They  
CC may also be used for inducing apoptosis and/or cell cycle arrest in  
CC a cell, particularly for treatment of cancer or other proliferative  
CC disorders such as psoriasis and restenosis.

XX  
SQ Sequence 9 AA;  
Query Match 80.6%; Score 25; DB 21; Length 9;

Best Local Similarity 100.0%; Pred. No. 9.3e+05;  
Matches 6; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 WXXWXF 6  
| | | | |  
Db 1 WXXWXF 6  
RESULT 43  
AAB20714  
ID AAB20714 standard; peptide; 11 AA.  
XX  
AC AAB20714;  
XX  
DT 20-DEC-2000 (first entry)  
XX  
DE Polymeric immunoglobulin receptor binding domain peptide SEQ ID NO:30.

XX  
KW Polymeric immunoglobulin receptor; pIgR; binding domain; diagnosis;  
KW identification; infection; cancer; asthma; antiasthmatic; cytostatic;  
KW antiinflammatory; antiinfectious; antidiarrhoeal; hepatotropic;  
KW virucide; vasotropic; anti-human immunodeficiency virus; antibacterial;  
KW mucosal epithelia; bronchitis; emphysema; cystic fibrosis; dysphagia;  
KW bronchiectasis; bronchiolitis; pulmonary oedema; viral tracheobronchitis;  
KW sleep apnea syndrome; infectious disease; neoplastic condition;  
KW Loffler's syndrome; kyphocloosis; peptic ulcer; diarrhoeal disease;  
KW ulcerative colitis; Crohn's disease; hepatitis; cirrhosis; haemorrhoid;  
KW systemic vasculitis; acquired immunodeficiency syndrome; gonorrhea;  
KW syphilis; chlamydia; antiulcer.

XX  
OS Homo sapiens.  
XX  
PN WO200047611-A2.

XX  
PD 17-AUG-2000.  
XX  
PF 11-FEB-2000; 2000WO-US03650.  
XX  
PR 12-FEB-1999; 99US-0119932.

XX  
PA (OKLA-) OKLAHOMA MEDICAL RES FOUND.  
PA (TEXA ) UNIV TEXAS SYSTEM.  
PA (DGIB-) DGI BIOTECHNOLOGIES.

XX  
PI Capra JD, White K, Hexham JM, Mandecki W;  
XX  
DR WPI; 2000-549134/50.

XX  
PT Novel polypeptides containing pIgR-binding domains used for targeting  
PT and transport to the mucosal epithelia, in the treatment of disorders  
PT accessible to the mucosal epithelia, e.g. asthma -

XX  
PS Claim 12; Fig 2; 139pp; English.

XX  
CC The present invention describes a 10-50 residue peptide (I) comprising  
CC a polymeric immunoglobulin receptor (pIgR)-binding domain. (I) can have  
CC antiasthmatic, antiinflammatory, antiinfectious, cytostatic, antiulcer,  
CC antidiarrhoeal, hepatotropic, virucide, vasotropic, anti-human  
CC immunodeficiency virus and antibacterial activities. (I) can be used  
CC for targeting and transport to the mucosal epithelium, for the  
CC prevention or treatment of diseases, ailments or conditions that are  
CC accessible to mucosal epithelia, including asthma, bronchitis, emphysema,  
CC cystic fibrosis, bronchiectasis, bronchiolitis, pulmonary oedema, viral  
CC tracheobronchitis, sleep apnea syndrome, infectious diseases, neoplastic  
CC conditions, Loffler's syndrome, kyphocloosis, dysphagia, peptic ulcers,  
CC diarrhoeal diseases, ulcerative colitis, Crohn's disease, hepatitis,  
CC cirrhosis, haemorrhoids, systemic vasculitis, acquired immunodeficiency  
CC syndrome, gonorrhea, syphilis and chlamydia. (I) can be attached to a  
CC detectable label for use in diagnostics. The present sequence represents  
CC a specifically claimed example of (I), derived from the human C-alpha-3  
CC domain amino acid sequence.

XX  
SQ Sequence 11 AA;



Query Match 80.6%; Score 25; DB 21; Length 11;  
Best Local Similarity 50.0%; Pred. No. 2.8e+02;  
Matches 3; Conservative 0; Mismatches 3; Indels 0; Gaps 0;

QY 1 WXXWXF 6  
| | |  
Db 3 WWSWLF 8.

RESULT 44  
AAW38112  
ID AAW38112 standard; Peptide; 13 AA.  
XX  
AC AAW38112;  
XX  
DT 23-APR-1998 (first entry)  
XX  
DE Dystrophin WW domain binding peptide 5.  
XX  
KW Peptide recognition unit; WW domain; cell signalling; growth regulation;  
KW cytoskeleton organisation; targeted drug screening; modulator;  
KW WW domain interaction; dystrophin.  
XX  
OS Synthetic.  
XX  
PN WO9737223-A1.  
XX  
PD 09-OCT-1997.  
XX  
PF 03-APR-1997; 97WO-US05547.  
XX  
PR 03-APR-1996; 96US-0630916.  
XX  
PA (CYTO-) CYTOGEN CORP.  
PA (UYNC-) UNIV NORTH CAROLINA.  
XX  
XX  
PI Fowlkes DM, Kay BK, Pirozzi G;  
DR WPI; 1997-503234/46.  
XX  
PT Identifying cell signalling and growth regulatory polypeptides by  
PT reaction with multivalent recognition complex - polypeptides are  
PT useful in targeted drug selection  
XX  
PS Example 2; Page 78; 220pp; English.  
XX  
CC Peptides AAW38108-13 function as recognition units of the dystrophin  
CC WW domain. They were identified from a random peptide phage display  
CC library using the dystrophin WW domain as a probe. The peptides were  
CC used as probes themselves to screen a lambda-EXlox mouse 16 day embryo  
CC cDNA expression library. In this way, cDNA clones expressing proteins,  
CC containing WW domains, capable of binding to these peptides are  
CC identified. The WW domain is a small functional domain found in a large  
CC number of proteins from a variety of species including humans, nematodes  
CC and yeast. Its name is derived from the observation that two tryptophan  
CC residues, one in the amino terminal portion of the WW domain and one in  
CC the carboxyl terminal portion, are conserved. Most proteins containing  
CC WW domains have a function involving cell signalling and growth  
CC regulation or the organisation of the cytoskeleton. Polypeptides  
CC containing a WW domain are identified by treating a multivalent  
CC recognition unit complex that has selective binding affinity for a WW  
CC domain, with many polypeptides and identifying those with selective  
CC affinity for the complex. Proteins containing WW domains are used for  
CC targeted drug screening, i.e. to identify potential modulators of  
CC specific WW domain interactions. The valency of the recognition unit is  
CC important in determining specificity of interaction with WW domains. In  
CC multivalent form specificity is relaxed, but not lost, so proteins  
CC containing WW domains similar, but not identical, to the sequence of  
CC the peptides' target WW can be detected, including new polypeptides.  
XX  
SQ Sequence 13 AA;

Query Match 80.6%; Score 25; DB 18; Length 13;  
Best Local Similarity 50.0%; Pred. No. 3.2e+02;  
Matches 3; Conservative 0; Mismatches 3; Indels 0; Gaps 0;

QY 1 WXXWXF 6  
| | |  
Db 5 WEEWEF 10

RESULT 45  
AAE07760  
ID AAE07760 standard; peptide; 14 AA.  
XX  
AC AAE07760;  
XX  
DT 06-NOV-2001 (first entry)  
XX  
DE Human HLA-DP restricted T cell epitope #4 of NY ESO-1 protein.  
XX  
KW Human; major histocompatibility complex; MHC; vaccine; metastasis;  
KW class II restricted T cell epitope; MHC-II epitope; cancer antigen;  
KW NY ESO-1 protein; CD4+ T lymphocyte; human leucocyte antigen; HLA;  
KW tumour-specific humoral-mediated immunity; cancer; cytostatic;  
KW immunotherapy.  
XX  
OS Homo sapiens.  
XX  
PN WO200155393-A2.  
XX  
PD 02-AUG-2001.  
XX  
PF 26-JAN-2001; 2001WO-US02765.  
XX  
PR 28-JAN-2000; 2000US-0179004.  
PR 29-SEP-2000; 2000US-0237107.  
XX  
XX (USSH ) US DEPT HEALTH & HUMAN SERVICES.  
PI Wang R, Rosenberg SA, Zeng G;  
XX WPI; 2001-496851/54.  
DR  
XX  
XX New NY-ESO cancer peptide or MHC class II restricted T cell epitopes,  
PT useful as immunogen and vaccine for inhibiting cancer in a mammal or as  
PT protection from metastasis -  
XX  
PS Claim 65; Page 82; 134pp; English.  
XX  
CC The invention relates to the identification and isolation of major  
CC histocompatibility (MHC) class II restricted T cell epitope (MHC-II  
CC epitope) derived from the cancer antigen, NY ESO-1. The MHC-II epitopes  
CC from NY ESO-1 are recognised by CD4+ T lymphocytes in an human leucocyte  
CC antigen (HLA) class II restricted manner, in particular HLA-DR or HLA-DP  
CC restricted. The products of the gene are promising candidates for  
CC immunotherapeutic strategies for the prevention, treatment and diagnosis  
CC of patients with cancer. The cancer epitopes are useful as immunogen and  
CC vaccine to inhibit or to prevent cancer in a mammal by eliciting CD4+ T  
CC lymphocytes resulting in protection of the recipient from development of  
CC cancer and protection from metastasis, or by inhibiting the growth of  
CC cells expressing the NY-ESO-1 gene product. The cancer peptides are also  
CC useful as diagnostic agent to detect the presence of cancer, to enhance  
CC the generation of antibody and/or CD8+ T cell responses against any  
CC given target antigen and/or hapten and to induce tumour-specific  
CC humoral-mediated immunity against cancer. The present sequence is human  
CC HLA-DP restricted T cell epitope of NY ESO-1 protein.  
XX  
SQ Sequence 14 AA;

Query Match 80.6%; Score 25; DB 22; Length 14;  
Best Local Similarity 50.0%; Pred. No. 3.3e+02;  
Matches 3; Conservative 0; Mismatches 3; Indels 0; Gaps 0;

QY 1 WXXWXF 6



Db           | | |  
              3 WITWCF 8

Search completed: December 12, 2003, 10:29:03  
Job time : 31.3 secs

GenCore version 5.1.6  
Copyright (c) 1993 - 2003 Compugen Ltd.

OM protein - protein search, using sw model

Run on: December 3, 2003, 11:48:05 ; Search time 26.3333 Seconds  
(without alignments)  
58.797 Million cell updates/sec

Title: US-09-912-414-11  
Perfect score: 38  
Sequence: 1 WXXWHF 6

Scoring table: BLOSUM62  
Gapop 10.0 , Gapext 0.5

Searched: 830525 seqs, 258052604 residues

Total number of hits satisfying chosen parameters: 3526

Minimum DB seq length: 0  
Maximum DB seq length: 15

Post-processing: Minimum Match 0%  
Maximum Match 100%  
Listing first 45 summaries

Database : SPTREMBL 23.\*

- 1: sp\_archaea.\*
- 2: sp\_bacteria.\*
- 3: sp\_fungi.\*
- 4: sp\_human.\*
- 5: sp\_invertebrate.\*
- 6: sp\_mammal.\*
- 7: sp\_mhc.\*
- 8: sp\_organelle.\*
- 9: sp\_phage.\*
- 10: sp\_plant.\*
- 11: sp\_rodent.\*
- 12: sp\_virus.\*
- 13: sp\_vertebrate.\*
- 14: sp\_unclassified.\*
- 15: sp\_rvirus.\*
- 16: sp\_bacteriap.\*
- 17: sp\_archaeap.\*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	DB ID	Description
1	22	57.9	8	13	P79940 xenopus lae
2	21	55.3	9	2	Q9R5M1 staphylococ
3	21	55.3	9	9	Q38366 bacterioph
4	20	52.6	9	8	Q8SHF0 chamaeleo n
5	20	52.6	12	7	O77919 pseudotroph
6	20	52.6	13	4	Q16406 homo sapien
7	20	52.6	15	2	Q53580 rhodobacter
8	19	50.0	8	8	Q94VF6 varanus job
9	19	50.0	10	13	Q9PRU9 sparus aura
10	19	50.0	14	8	Q9MT61 allium cepa
11	19	50.0	14	8	Q9MRV4 allium porr
12	19	50.0	14	8	Q9MRV1 allium sati
13	19	50.0	14	8	Q9MRT8 aloe vera (
14	19	50.0	14	8	Q8HGT1 gadus morhu
15	17	44.7	10	8	Q94VD2 varanus pan
16	17	44.7	13	10	Q8LPV3 deschampsia

17	44.7	14	6	Q9TQZ1	Q9tqz1 bos taurus
18	44.7	14	11	Q9R1G8	Q9rlg8 rattus norv
19	42.1	8	8	Q94VCL	Q94vcl varanus rud
20	42.1	8	8	Q9TD02	Q9td02 terranatos
21	42.1	8	8	Q9T4Y2	Q9t4y2 asterina pe
22	42.1	9	8	Q9T688	Q9t688 gecko gecko
23	42.1	10	2	Q47561	Q47561 escherichia
24	42.1	10	8	Q9T8K7	Q9t8k7 liolaemus m
25	42.1	10	8	Q9T8N1	Q9t8n1 liolaemus p
26	42.1	10	8	Q79903	Q79903 oplurus cuv
27	42.1	10	8	Q8W969	Q8w969 anolis orto
28	42.1	10	8	Q8WDH8	Q8wdh8 anolis mest
29	42.1	10	8	Q9T8T6	Q9t8t6 liolaemus m
30	42.1	10	8	Q9T8L3	Q9t8l3 liolaemus l
31	42.1	10	8	P92616	P92616 aspidosceli
32	42.1	10	8	Q9T8G8	Q9t8g8 liolaemus c
33	42.1	10	8	Q9S8K9	Q9s8k9 rana boylii
34	42.1	10	8	Q9TFU9	Q9tfu9 teratoscinc
35	42.1	10	8	Q9T8X7	Q9t8x7 phymaturus
36	42.1	10	8	Q9S8L2	Q9s8l2 rana tempor
37	42.1	10	8	Q79885	Q79885 anolis pate
38	42.1	10	8	Q9T8Q5	Q9t8q5 liolaemus l
39	42.1	10	8	P92654	P92654 euprepis au
40	42.1	10	8	Q9T8L0	Q9t8l0 liolaemus o
41	42.1	10	8	Q9T8W8	Q9t8w8 liolaemus b
42	42.1	10	8	Q9T8R4	Q9t8r4 liolaemus p
43	42.1	10	8	Q9T8M8	Q9t8m8 liolaemus m
44	42.1	10	8	Q9T8S1	Q9t8s1 liolaemus l
45	42.1	10	8	Q9T8S4	Q9t8s4 liolaemus c

ALIGNMENTS

RESULT 1  
P79940  
ID P79940 PRELIMINARY; PRT; 8 AA.  
AC P79940;  
DT 01-MAY-1997 (TrEMBLrel. 03, Created)  
DT 01-MAY-1997 (TrEMBLrel. 03, Last sequence update)  
DT 01-DEC-2001 (TrEMBLrel. 19, Last annotation update)  
DE XMeisl-4 protein (Fragment).  
OS Xenopus laevis (African clawed frog).  
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
OC Amphibia; Batrachia; Anura; Mesobatrachia; Pipidae;  
OC Xenopodinae; Xenopus.  
OX NCBI\_TaxID=8355;  
RN [1]  
RP SEQUENCE FROM N.A.  
RX MEDLINE=97202105; PubMed=9049632;  
RA Steelman S., Moskow J.J., Muzynski K., North C., Druck T.,  
RA Montgomery J.C., Huebner K., Daar I.O., Buchberg A.M.;  
RT "Identification of a conserved family of Meisl-related homeobox genes."  
RL Genome Res. 7:142-156(1997).  
DR EMBL; U68389; AAB19199.1; -.  
DR TRANSFAC; T03410; -.  
FT NON TER 1 1  
SQ SEQUENCE 8 AA; 1187 MW; 278B51F37B11F40B CRC64;  
Query Match 57.9%; Score 22; DB 13; Length 8;  
Best Local Similarity 66.7%; Pred. No. 8.3e+05;  
Matches 2; Conservative 1; Mismatches 0; Indels 0; Gaps 0;  
QY 4 WHF 6  
Db ||:  
5 WHY 7

RESULT 2  
Q9R5M1  
ID Q9R5M1 PRELIMINARY; PRT; 9 AA.  
AC Q9R5M1;

DT 01-MAY-2000 (TrEMBLrel. 13, Created)  
DT 01-MAY-2000 (TrEMBLrel. 13, Last sequence update)  
DT 01-JUN-2002 (TrEMBLrel. 21, Last annotation update)  
DE 66 kDa cell surface adhesin for heparan sulfate (Fragment).  
OS Staphylococcus aureus.  
OC Bacteria; Firmicutes; Bacillales; Staphylococcus.  
OX NCBI\_TaxID=1280;  
RN [1]  
RP SEQUENCE.  
RX MEDLINE=92176005; PubMed=1541563;  
RA Liang O.D., Ascencio F.; Fransson L.A., Wadstrom T.;  
RT "Binding of heparan sulfate to Staphylococcus aureus.";  
RL Infect. Immun. 60:899-906(1992).  
FT NON\_TER 1  
FT NON\_TER 9  
SQ SEQUENCE 9 AA; 990 MW; 2289DDD7337861B3 CRC64;

Query Match 55.3%; Score 21; DB 2; Length 9;  
Best Local Similarity 50.0%; Pred. No. 8.3e+05;  
Matches 2; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

Qy 1 WXXW 4  
| |  
Db 2 WTCW 5

RESULT 3  
Q38366  
ID Q38366 PRELIMINARY; PRT; 9 AA.  
AC Q38366;  
DT 01-NOV-1996 (TrEMBLrel. 01, Created)  
DT 01-NOV-1996 (TrEMBLrel. 01, Last sequence update)  
DT 01-DEC-2001 (TrEMBLrel. 19, Last annotation update)  
DE E gene product (Fragment).  
OS Bacteriophage phi-X174.  
OC Viruses; ssDNA viruses; Microviridae; Microvirus.  
OX NCBI\_TaxID=10847;  
RN [1]  
RP SEQUENCE FROM N.A.  
RX MEDLINE=88118956; PubMed=2963134;  
RA Buckley K.J., Hayashi M.;  
RT "Role of premature translational termination in the regulation of  
RT expression of the phiX174 lysis gene.";  
RL J. Mol. Biol. 198:599-607(1987).  
DR EMBL; X07809; CAA30668.1; -.  
FT NON\_TER 9  
FT NON\_TER 9  
SQ SEQUENCE 9 AA; 1207 MW; C093B37731B36412 CRC64;

Query Match 55.3%; Score 21; DB 9; Length 9;  
Best Local Similarity 50.0%; Pred. No. 8.3e+05;  
Matches 2; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

Qy 1 WXXW 4  
| |  
Db 4 WTLW 7

RESULT 4  
Q8SHF0  
ID Q8SHF0 PRELIMINARY; PRT; 9 AA.  
AC Q8SHF0;  
DT 01-JUN-2002 (TrEMBLrel. 21, Created)  
DT 01-JUN-2002 (TrEMBLrel. 21, Last sequence update)  
DT 01-JUN-2002 (TrEMBLrel. 21, Last annotation update)  
DE Cytochrome c oxidase subunit I (Fragment).  
GN COI.  
OS Chamaeleo namaquensis.  
OC Mitochondrion.  
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
OC Lepidosauria; Squamata; Iguania; Acrodonta; Chamaeleonidae; Chamaeleo.  
OX NCBI\_TaxID=179917;  
RN [1]  
RP SEQUENCE FROM N.A.

RA Townsend T.M., Larson A.L.;  
RT "Molecular Phylogenetics and Mitochondrial Genomic Evolution in the  
RT Chamaeleonidae (Reptilia, Squamata).";  
RL Submitted (NOV-2001) to the EMBL/GenBank/DBJ databases.  
DR EMBL; AF448757; AAL90553.1; -.  
KW Mitochondrion.  
FT NON\_TER 9  
FT NON\_TER 9  
SQ SEQUENCE 9 AA; 1205 MW; 358CB72733640733 CRC64;

Query Match 52.6%; Score 20; DB 8; Length 9;  
Best Local Similarity 50.0%; Pred. No. 8.3e+05;  
Matches 2; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

Qy 1 WXXW 4  
| |  
Db 2 WLRW 5

RESULT 5  
O77919  
ID O77919 PRELIMINARY; PRT; 12 AA.  
AC O77919;  
DT 01-NOV-1998 (TrEMBLrel. 08, Created)  
DT 01-NOV-1998 (TrEMBLrel. 08, Last sequence update)  
DT 01-DEC-2001 (TrEMBLrel. 19, Last annotation update)  
DE MHC class II B locus 4 (Fragment).  
OS Pseudotropheus sp. 'pseudotropheus tropheops complex'.  
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
OC Actinopterygii; Neopterygii; Teleostei; Euteleostei; Neoteleostei;  
OC Acanthomorpha; Acanthopterygii; Percomorpha; Perciformes; Labroidae;  
OC Cichlidae; Pseudotropheus.  
OX NCBI\_TaxID=51796;  
RN [1]  
RP SEQUENCE FROM N.A.  
RX MEDLINE=98315113; PubMed=9649539;  
RA Malaga-Trillo E., Zaleska-Rutczynska Z., McAndrew B., Vincek V.,  
RA Figueroa F., Sultmann H., Klein J.;  
RT "Linkage relationships and haplotype polymorphism among cichlid mhc  
RT class II B loci.";  
RL Genetics 149:1527-1537(1998).  
DR EMBL; AF050032; AAC41371.1; -.  
FT NON\_TER 1  
FT NON\_TER 12  
FT NON\_TER 12  
SQ SEQUENCE 12 AA; 1529 MW; 6C2ABFACD5A5B734 CRC64;

Query Match 52.6%; Score 20; DB 7; Length 12;  
Best Local Similarity 50.0%; Pred. No. 2.3e+03;  
Matches 2; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

Qy 1 WXXW 4  
| |  
Db 1 WDFW 4

RESULT 6  
Q16406  
ID Q16406 PRELIMINARY; PRT; 13 AA.  
AC Q16406;  
DT 01-NOV-1996 (TrEMBLrel. 01, Created)  
DT 01-NOV-1996 (TrEMBLrel. 01, Last sequence update)  
DT 01-MAY-1999 (TrEMBLrel. 10, Last annotation update)  
DE GHRH-R protein (Fragment).  
GN GHRH-R.  
OS Homo sapiens (Human).  
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
OC Mammalia; Euthera; Primates; Catarrhini; Homnidae; Homo.  
OX NCBI\_TaxID=9606;  
RN [1]  
RP SEQUENCE FROM N.A.  
RX MEDLINE=96001284; PubMed=7559877;  
RA Hashimoto K., Koga M., Motomura T., Kasayama S., Kouhara H.,  
RA Ohnishi T., Arita N., Hayakawa T., Sato B., Kishimoto T.;  
RT "Identification of alternatively spliced messenger ribonucleic acid

RT encoding truncated growth hormone-releasing hormone receptor in human  
RT pituitary adenomas."  
RT J. Clin. Endocrinol. Metab. 80:2933-2939(1995).  
DR EMBL; S79912; AAD14318.1; --  
FT NON TER 1  
SQ SEQUENCE 13 AA; 1612 MW; CE19D7D255D66362 CRC64;

Query Match 52.6%; Score 20; DB 4; Length 13;  
Best Local Similarity 50.0%; Pred. No. 2.5e+03;  
Matches 2; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

QY 1 WXXW 4  
| |  
Db 7 WGYW 10

## RESULT 7

Q53580 PRELIMINARY; PRT; 15 AA.  
AC Q53580;  
DT 01-NOV-1996 (TrEMBLrel. 01, Created)  
DT 01-NOV-1996 (TrEMBLrel. 01, Last sequence update)  
DT 01-DEC-2001 (TrEMBLrel. 19, Last annotation update)  
DE Light-harvesting complex I alpha polypeptide (Fragment).  
GN PUFA.  
OS Rhodobacter capsulatus (Rhodospseudomonas capsulata).  
OC Bacteria; Proteobacteria; Alphaproteobacteria; Rhodobacterales;  
OC Rhodobacteraceae; Rhodobacter.  
OX NCBI\_TaxID=1061;  
RN [1]  
RP SEQUENCE FROM N.A.  
RX MEDLINE=92234963; PubMed=1569029;  
RA Richter P.; Brand M.; Drews G.;  
RT "Characterization of LHI- and LHI+ Rhodobacter capsulatus pufa  
mutants.";  
RL J. Bacteriol. 174:3030-3041(1992).  
DR EMBL; S97552; AAC60406.1; --  
FT NON TER 15  
SQ SEQUENCE 15 AA; 2054 MW; 3561FE413591D31A CRC64;

Query Match 52.6%; Score 20; DB 2; Length 15;  
Best Local Similarity 50.0%; Pred. No. 2.8e+03;  
Matches 2; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

QY 1 WXXW 4  
| |  
Db 8 WKIW 11

## RESULT 8

Q94VF6 PRELIMINARY; PRT; 8 AA.  
AC Q94VF6;  
DT 01-DEC-2001 (TrEMBLrel. 19, Created)  
DT 01-DEC-2001 (TrEMBLrel. 19, Last sequence update)  
DT 01-DEC-2001 (TrEMBLrel. 19, Last annotation update)  
DE Cytochrome c oxidase subunit I (Fragment).  
GN COI.  
OS Varanus jobiensis.  
OG Mitochondrion.  
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
OC Lepidosauria; Squamata; Scleroglossa; Anguimorpha; Varanidae; Varanus.  
OX NCBI\_TaxID=169843;  
RN [1]  
RP SEQUENCE FROM N.A.  
RA Ast J.C.;  
RT "Mitochondrial DNA evidence and evolution in Varanoidea (Squamata).";  
RL Cladistics 17:0-0(2001).  
DR EMBL; AF407507; AAL10075.1; --  
KW Mitochondrion.  
FT NON TER 8  
SQ SEQUENCE 8 AA; 1144 MW; EFD729DB436411A6 CRC64;

Query Match 50.0%; Score 19; DB 8; Length 8;  
Best Local Similarity 66.7%; Pred. No. 8.3e+05;  
Matches 2; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

QY 4 WHF 6  
|:|  
Db 4 WYF 6

## RESULT 9

Q9PRU9 PRELIMINARY; PRT; 10 AA.  
AC Q9PRU9;  
DT 01-MAY-2000 (TrEMBLrel. 13, Created)  
DT 01-MAY-2000 (TrEMBLrel. 13, Last sequence update)  
DT 01-MAY-2000 (TrEMBLrel. 13, Last annotation update)  
DE Gonadotropin-releasing hormone, SGNRH-I.  
OS Sparus aurata (Gilthead sea bream).  
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
OC Actinopterygii; Neopterygii; Teleostei; Euteleostei; Neoteleostei;  
OC Acanthomorpha; Acanthopterygii; Percomorpha; Perciformes; Percoidae;  
OC Sparidae; Sparus.  
OX NCBI\_TaxID=8175;  
RN [1]  
RP SEQUENCE.  
RX MEDLINE=95083645; PubMed=7991588;  
RA Powell J.F.; Zohar Y.; Elizur A.; Park M.; Fischer W.H.; Craig A.G.;  
RA Rivier J.E.; Lovejoy D.A.; Sherwood N.M.;  
RT "Three forms of gonadotropin-releasing hormone characterized from  
brains of one species.";  
RL Proc. Natl. Acad. Sci. U.S.A. 91:12081-12085(1994).  
SQ SEQUENCE 10 AA; 1132 MW; 81566865AB587735 CRC64;

Query Match 50.0%; Score 19; DB 13; Length 10;  
Best Local Similarity 100.0%; Pred. No. 2.8e+03;  
Matches 2; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 4 WH 5  
||  
Db 8 WH 9

## RESULT 10

Q9MT61 PRELIMINARY; PRT; 14 AA.  
AC Q9MT61;  
DT 01-OCT-2000 (TrEMBLrel. 15, Created)  
DT 01-OCT-2000 (TrEMBLrel. 15, Last sequence update)  
DT 01-OCT-2000 (TrEMBLrel. 15, Last annotation update)  
DE PSI 9 kDa protein (Fragment).  
GN PSAC.  
OS Allium cepa (Onion).  
OG Chloroplast.  
OC Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;  
OC Spermatophyta; Magnoliophyta; Liliopsida; Asparagales; Alliaceae;  
OC Allium.  
OX NCBI\_TaxID=4679;  
RN [1]  
RP SEQUENCE FROM N.A.  
RC TISSUE=Leaf;  
RA Lopez-Serrano M.; del Campo E.M.; Sabater B.; Martin M.;  
RT "Conservation of the start codon by editing in nhhd-encoded  
transcripts is not restricted to dicotyledonous plants.";  
RL Submitted (JUN-2000) to the EMBL/GenBank/DBJ databases.  
DR EMBL; AJ278350; CAB96183.1; --  
KW Chloroplast.  
FT NON TER 1  
SQ SEQUENCE 14 AA; 1744 MW; 8F14FD03E3B7D911 CRC64;

Query Match 50.0%; Score 19; DB 8; Length 14;  
Best Local Similarity 100.0%; Pred. No. 3.8e+03;  
Matches 2; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 4 WH 5  
||  
Db 3 WH 4

## RESULT 11

Q9MRV4 ID Q9MRV4 PRELIMINARY; PRT; 14 AA.  
AC Q9MRV4;  
DT 01-OCT-2000 (TrEMBLrel. 15, Created)  
DT 01-OCT-2000 (TrEMBLrel. 15, Last sequence update)  
DT 01-OCT-2000 (TrEMBLrel. 15, Last annotation update)  
DE PSI 9 kDa protein (Fragment).  
GN PSAC.  
OS Allium porrum (Leek).  
OG Chloroplast.  
OC Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;  
OC Spermatophyta; Magnoliophyta; Liliopsida; Asparagales; Alliaceae;  
OC Allium.  
OX NCBI\_TaxID=4681;  
RN [1]  
RP SEQUENCE FROM N.A.  
RC TISSUE=Leaf;  
RA Lopez-Serrano M., del Campo E.M., Sabater B., Martin M.;  
RT "Conservation of the start codon by editing in ndhD-encoded  
transcripts is not restricted to dicotyledonous plants.";  
RL Submitted (JUN-2000) to the EMBL/GenBank/DBJ databases.  
DR EMBL; AJ278352; CAB96185.1; -.  
KW Chloroplast.  
FT NON\_TER 1  
SQ SEQUENCE 14 AA; 1744 MW; 8F14FD03E3B7D911 CRC64;

Query Match 50.0%; Score 19; DB 8; Length 14;  
Best Local Similarity 100.0%; Pred. No. 3.8e+03;  
Matches 2; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 4 WH 5  
||  
Db 3 WH 4

## RESULT 12

Q9MRV1 ID Q9MRV1 PRELIMINARY; PRT; 14 AA.  
AC Q9MRV1;  
DT 01-OCT-2000 (TrEMBLrel. 15, Created)  
DT 01-OCT-2000 (TrEMBLrel. 15, Last sequence update)  
DT 01-OCT-2000 (TrEMBLrel. 15, Last annotation update)  
DE PSI 9 kDa protein (Fragment).  
GN PSAC.  
OS Allium sativum (Garlic).  
OG Chloroplast.  
OC Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;  
OC Spermatophyta; Magnoliophyta; Liliopsida; Asparagales; Alliaceae;  
OC Allium.  
OX NCBI\_TaxID=4682;  
RN [1]  
RP SEQUENCE FROM N.A.  
RC TISSUE=Leaf;  
RA Lopez-Serrano M., del Campo E.M., Sabater B., Martin M.;  
RT "Conservation of the start codon by editing in ndhD-encoded  
transcripts is not restricted to dicotyledonous plants.";  
RL Submitted (JUN-2000) to the EMBL/GenBank/DBJ databases.  
DR EMBL; AJ278351; CAB96187.1; -.  
KW Chloroplast.  
FT NON\_TER 1  
SQ SEQUENCE 14 AA; 1744 MW; 8F14FD03E3B7D911 CRC64;

Query Match 50.0%; Score 19; DB 8; Length 14;  
Best Local Similarity 100.0%; Pred. No. 3.8e+03;  
Matches 2; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 4 WH 5  
||

Db 3 WH 4  
||

## RESULT 13

Q9MRT8 ID Q9MRT8 PRELIMINARY; PRT; 14 AA.  
AC Q9MRT8;  
DT 01-OCT-2000 (TrEMBLrel. 15, Created)  
DT 01-OCT-2000 (TrEMBLrel. 15, Last sequence update)  
DT 01-DEC-2001 (TrEMBLrel. 19, Last annotation update)  
DE PSI 9 kDa protein (Fragment).  
GN PSAC.  
OS Aloe vera (Aloe) (Aloe barbadensis).  
OG Chloroplast.  
OC Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;  
OC Spermatophyta; Magnoliophyta; Liliopsida; Asparagales; Asphodelaceae;  
OC Aloe.  
OX NCBI\_TaxID=34199;  
RN [1]  
RP SEQUENCE FROM N.A.  
RC TISSUE=Leaf;  
RA Lopez-Serrano M., del Campo E.M., Sabater B., Martin M.;  
RT "Conservation of the start codon by editing in ndhD-encoded  
transcripts is not restricted to dicotyledonous plants.";  
RL Submitted (JUN-2000) to the EMBL/GenBank/DBJ databases.  
DR EMBL; AJ278353; CAB96192.1; -.  
KW Chloroplast.  
FT NON\_TER 1  
SQ SEQUENCE 14 AA; 1744 MW; 8F14FD03E3B7D911 CRC64;

Query Match 50.0%; Score 19; DB 8; Length 14;  
Best Local Similarity 100.0%; Pred. No. 3.8e+03;  
Matches 2; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 4 WH 5  
||  
Db 3 WH 4

## RESULT 14

Q8HGT1 ID Q8HGT1 PRELIMINARY; PRT; 14 AA.  
AC Q8HGT1;  
DT 01-MAR-2003 (TrEMBLrel. 23, Created)  
DT 01-MAR-2003 (TrEMBLrel. 23, Last sequence update)  
DT 01-MAR-2003 (TrEMBLrel. 23, Last annotation update)  
DE AtPase 8 (Fragment).  
OS Gadus morhua (Atlantic cod).  
OG Mitochondrion.  
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
OC Actinopterygii; Neopterygii; Teleostei; Euteleostei; Neoteleostei;  
OC Acanthomorpha; Paracanthopterygii; Gadiformes; Gadidae; Gadus.  
OX NCBI\_TaxID=8049;  
RN [1]  
RP SEQUENCE FROM N.A.  
RC STRAIN=ATPRK3;  
RA Taylor M.I., Fox C., Rico I., Rico C.;  
RT "Species-specific TagMan probes for simultaneous identification of  
(Gadus morhua L.), haddock (Melanogrammus aeglefinus L.) and whiting  
(Merlangius merlangus L.).";  
RL Mol. Ecol. Notes 2:599-601 (2002).  
DR EMBL; AF526615; AAN85062.1; -.  
KW Mitochondrion.  
FT NON\_TER 1  
SQ SEQUENCE 14 AA; 1753 MW; D4AF852330085E6D CRC64;

Query Match 50.0%; Score 19; DB 8; Length 14;  
Best Local Similarity 100.0%; Pred. No. 3.8e+03;  
Matches 2; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 4 WH 5  
||



Db 13 WH 14

RESULT 15

Q94VD2 Q94VD2 PRELIMINARY; PRT; 10 AA.  
AC Q94VD2;  
DT 01-DEC-2001 (TReMBLrel. 19, Created)  
DT 01-DEC-2001 (TReMBLrel. 19, Last sequence update)  
DT 01-DEC-2001 (TReMBLrel. 19, Last annotation update)  
DE Cytochrome c oxidase subunit I (Fragment).  
GN COI.  
OS Varanus panoptes panoptes.  
OG Mitochondrion.  
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
OC Lepidosauria; Squamata; Scleroglossa; Anguimorpha; Varanidae; Varanus.  
OX NCBI\_TaxID=169849;  
RN [1]  
RP SEQUENCE FROM N.A.  
RA Ast J.C.;  
RT "Mitochondrial DNA evidence and evolution in Varanoidea (Squamata).";  
RL Cladistics 17:0-0(2001).  
DR EMBL; AF407516; AAL10102.1; -.  
KW Mitochondrion.  
FT NON TER 10 10  
SQ SEQUENCE 10 AA; 1299 MW; 5DEE80D4136411A7 CRC64;

Query Match 44.7%; Score 17; DB 8; Length 10;  
Best Local Similarity 66.7%; Pred. No. 6e+03;  
Matches 2; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

QY 4 WHF 6  
Db -6 WRF 8

Search completed: December 3, 2003, 11:53:25  
Job time : 27.3333 secs

GenCore version 5.1.6  
Copyright (c) 1993 - 2003 Compugen Ltd.

OM protein - protein search, using sw model

Run on: December 3, 2003, 11:44:20 ; Search time 33.6667 Seconds  
(without alignments)  
28.288 Million cell updates/sec

Title: US-09-912-414-11  
Perfect score: 38  
Sequence: 1 WXXWHF 6

Scoring table: BLOSUM62  
Gapop 10.0 , Gapext 0.5

Searched: 1107863 seqs, 158726573 residues

Total number of hits satisfying chosen parameters: 350435

Minimum DB seq length: 0  
Maximum DB seq length: 15

Post-processing: Minimum Match 0%  
Maximum Match 100%  
Listing first 45 summaries

Database : A Geneseq 19Jun03:\*  
1: /SIDS1/gcgdata/geneseq/geneseq-emb1/AA1980.DAT:\*  
2: /SIDS1/gcgdata/geneseq/geneseq-emb1/AA1981.DAT:\*  
3: /SIDS1/gcgdata/geneseq/geneseq-emb1/AA1982.DAT:\*  
4: /SIDS1/gcgdata/geneseq/geneseq-emb1/AA1983.DAT:\*  
5: /SIDS1/gcgdata/geneseq/geneseq-emb1/AA1984.DAT:\*  
6: /SIDS1/gcgdata/geneseq/geneseq-emb1/AA1985.DAT:\*  
7: /SIDS1/gcgdata/geneseq/geneseq-emb1/AA1986.DAT:\*  
8: /SIDS1/gcgdata/geneseq/geneseq-emb1/AA1987.DAT:\*  
9: /SIDS1/gcgdata/geneseq/geneseq-emb1/AA1988.DAT:\*  
10: /SIDS1/gcgdata/geneseq/geneseq-emb1/AA1989.DAT:\*  
11: /SIDS1/gcgdata/geneseq/geneseq-emb1/AA1990.DAT:\*  
12: /SIDS1/gcgdata/geneseq/geneseq-emb1/AA1991.DAT:\*  
13: /SIDS1/gcgdata/geneseq/geneseq-emb1/AA1992.DAT:\*  
14: /SIDS1/gcgdata/geneseq/geneseq-emb1/AA1993.DAT:\*  
15: /SIDS1/gcgdata/geneseq/geneseq-emb1/AA1994.DAT:\*  
16: /SIDS1/gcgdata/geneseq/geneseq-emb1/AA1995.DAT:\*  
17: /SIDS1/gcgdata/geneseq/geneseq-emb1/AA1996.DAT:\*  
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21: /SIDS1/gcgdata/geneseq/geneseq-emb1/AA2000.DAT:\*  
22: /SIDS1/gcgdata/geneseq/geneseq-emb1/AA2001.DAT:\*  
23: /SIDS1/gcgdata/geneseq/geneseq-emb1/AA2002.DAT:\*  
24: /SIDS1/gcgdata/geneseq/geneseq-emb1/AA2003.DAT:\*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	ID	Description
1	35	92.1	6	21	AA01505
2	35	92.1	6	21	AA01506
3	34	89.5	6	21	AA01492
4	34	89.5	6	21	AA01499
5	29	76.3	6	24	ABR45593
6	29	76.3	6	24	ABR45594
7	29	76.3	14	22	AA00214
8	28	73.7	6	18	AAW28912
9	28	73.7	6	18	AA093770
					Peptide which bind
					Peptide which bind
					Peptide which bind
					Peptide which bind
					Staphylococcus aur
					Staphylococcus aur
					Human angiotensin
					Opioid peptide. S
					New peptide which

10	28	73.7	6	20	AA23019	Opioid peptide whi
11	28	73.7	6	21	AA01509	Peptide which bind
12	28	73.7	6	24	ABR45591	Staphylococcus aur
13	28	73.7	6	24	ABR45592	Staphylococcus aur
14	28	73.7	7	20	AA01258	US5851813 peptide
15	28	73.7	7	22	AA049729	Peptide SEQ ID 40
16	28	73.7	8	15	AA060429	Antiproliferative
17	28	73.7	8	15	AA060444	Antiproliferative
18	28	73.7	8	16	AA083499	Zif268 mutagenised
19	28	73.7	8	20	AA01261	US5851813 peptide
20	28	73.7	8	20	AA084388	Finger 3 binding s
21	28	73.7	12	21	AA088108	Fluorescein bindin
22	28	73.7	12	21	AA088160	Fluorescein bindin
23	28	73.7	12	22	AA060032	Internalising pept
24	27	71.1	6	19	AA083884	Peptide specific a
25	27	71.1	7	22	AA045777	H11 binding site c
26	27	71.1	8	23	AB0493	Hominidae LDL rece
27	27	71.1	10	18	AA032766	Human platelet gly
28	27	71.1	11	22	AA012188	Polyglutamine-glut
29	27	71.1	13	23	AB046201	Human BLYS binding
30	26	68.4	6	24	ABR45313	Staphylococcus aur
31	26	68.4	6	24	ABR45314	Staphylococcus aur
32	26	68.4	6	24	ABR47161	Staphylococcus aur
33	26	68.4	6	24	ABR47162	Staphylococcus aur
34	26	68.4	9	23	AA026775	Fibrin binding pep
35	26	68.4	9	23	AA093672	Granulocyte-colony
36	26	68.4	12	21	AA088104	Oregon green 514 b
37	26	68.4	12	21	AA088154	Oregon green 514 b
38	26	68.4	13	18	AA038112	Dystrophin ww doma
39	26	68.4	15	20	AA030351	Epitope derived fr
40	26	68.4	15	23	AA026759	Fibrin binding pep
41	26	68.4	15	23	AA019239	Streptococcus pneu
42	25.5	67.1	15	23	AB057721	Human nucleotide e
43	25	65.8	6	13	AA024966	Phe-Arg contg. ant
44	25	65.8	6	15	AA057391	Peptide for treati
45	25	65.8	6	21	AA01493	Peptide which bind

ALIGNMENTS

RESULT 1  
AA01505  
ID AA01505 standard; peptide; 6 AA.  
XX  
AC AA01505;  
XX  
DT 08-NOV-2000 (first entry)  
XX  
DE Peptide which binds to transcription factor E2F-1 DNA binding domain.  
XX  
KW DNA binding; transcription factor; E2F; E2F-1; cell cycle; DP-1;  
KW activation; transcription; apoptosis; proliferative disorder;  
KW psoriasis; restenosis.  
XX  
OS Synthetic.  
XX  
PN WO200044771-A1.  
XX  
PD 03-AUG-2000.  
XX  
PF 26-JAN-2000; 2000WO-GB00227.  
XX  
PR 26-JAN-1999; 99GB-0001710.  
XX  
PA (PROL-) PROLIFIX LTD.  
XX  
PI Mueller R, Kontermann RE, Montigiani S;  
XX  
DR WPI; 2000-532806/48.  
XX  
PT Peptides binding to the DNA binding domain of transcription factor E2F and inhibiting cell cycle progression, useful for the treatment of

PT Cancer

PS Example; Page 26; 42pp; English.

XX

CC Peptides which bind to the DNA binding domain of transcription factor E2F and inhibit cell cycle progression may be useful as research agents to investigate the interaction between E2F and DP-1, or the activation of transcription by E2F-1/DP-1 heterodimers. They may also be used for inducing apoptosis and/or cell cycle arrest in a cell, particularly for treatment of cancer or other proliferative disorders such as psoriasis and restenosis.

XX

SQ Sequence 6 AA;

Query Match 92.1%; Score 35; DB 21; Length 6;

Best Local Similarity 66.7%; Pred. No. 9.3e+05;

Matches 4; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

OY 1 WXXWHF 6

Db 1 WARWHF 6

RESULT 2

AAB01506

ID AAB01506 standard; peptide; 6 AA.

XX

AC AAB01506;

XX

DT 08-NOV-2000 (first entry)

XX

DE Peptide which binds to transcription factor E2F-1 DNA binding domain.

XX

KW DNA binding; transcription factor; E2F; E2F-1; cell cycle; DP-1; activation; transcription; apoptosis; proliferative disorder; psoriasis; restenosis.

KW

XX

OS Synthetic.

XX

PN WO200044771-A1.

XX

PD 03-AUG-2000.

XX

PF 26-JAN-2000; 2000WO-GB00227.

XX

PR 26-JAN-1999; 99GB-0001710.

XX

PA (PROL-) PROLIFIX LTD.

XX

PI Mueller R, Kontermann RE, Montigiani S;

XX

DR WPI; 2000-532806/48.

XX

PD 03-AUG-2000.

XX

PF 26-JAN-2000; 2000WO-GB00227.

XX

PR 26-JAN-1999; 99GB-0001710.

XX

PA (PROL-) PROLIFIX LTD.

XX

PI Mueller R, Kontermann RE, Montigiani S;

XX

DR WPI; 2000-532806/48.

XX

PT Peptides binding to the DNA binding domain of transcription factor E2F and inhibiting cell cycle progression, useful for the treatment of cancer

PT

XX

PS Example; Page 26; 42pp; English.

XX

CC Peptides which bind to the DNA binding domain of transcription factor E2F and inhibit cell cycle progression may be useful as research agents to investigate the interaction between E2F and DP-1, or the activation of transcription by E2F-1/DP-1 heterodimers. They may also be used for inducing apoptosis and/or cell cycle arrest in a cell, particularly for treatment of cancer or other proliferative disorders such as psoriasis and restenosis.

XX

SQ Sequence 6 AA;

Query Match 92.1%; Score 35; DB 21; Length 6;

Best Local Similarity 66.7%; Pred. No. 9.3e+05;

Matches 4; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

OY 1 WXXWHF 6

Db 1 WARWHF 6

RESULT 2

AAB01506

ID AAB01506 standard; peptide; 6 AA.

XX

AC AAB01506;

XX

DT 08-NOV-2000 (first entry)

XX

DE Peptide which binds to transcription factor E2F-1 DNA binding domain.

XX

KW DNA binding; transcription factor; E2F; E2F-1; cell cycle; DP-1; activation; transcription; apoptosis; proliferative disorder; psoriasis; restenosis.

KW

XX

OS Synthetic.

XX

PN WO200044771-A1.

XX

PD 03-AUG-2000.

XX

PF 26-JAN-2000; 2000WO-GB00227.

XX

PR 26-JAN-1999; 99GB-0001710.

XX

PA (PROL-) PROLIFIX LTD.

XX

PI Mueller R, Kontermann RE, Montigiani S;

XX

DR WPI; 2000-532806/48.

XX

PT Peptides binding to the DNA binding domain of transcription factor E2F and inhibiting cell cycle progression, useful for the treatment of cancer

PT

XX

PS Example; Page 26; 42pp; English.

XX

CC Peptides which bind to the DNA binding domain of transcription factor E2F and inhibit cell cycle progression may be useful as research agents to investigate the interaction between E2F and DP-1, or the activation of transcription by E2F-1/DP-1 heterodimers. They may also be used for inducing apoptosis and/or cell cycle arrest in a cell, particularly for treatment of cancer or other proliferative disorders such as psoriasis and restenosis.

XX

SQ Sequence 6 AA;

Query Match 92.1%; Score 35; DB 21; Length 6;

Best Local Similarity 66.7%; Pred. No. 9.3e+05;

Matches 4; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

OY 1 WXXWHF 6

Db 1 WARWHF 6

RESULT 2

AAB01506

ID AAB01506 standard; peptide; 6 AA.

XX

AC AAB01506;

XX

DT 08-NOV-2000 (first entry)

XX

DE Peptide which binds to transcription factor E2F-1 DNA binding domain.

XX

KW DNA binding; transcription factor; E2F; E2F-1; cell cycle; DP-1; activation; transcription; apoptosis; proliferative disorder; psoriasis; restenosis.

KW

OY 1 WXXWHF 6

Db 1 WVAWHF 6

RESULT 3

AAB01492

ID AAB01492 standard; peptide; 6 AA.

XX

AC AAB01492;

XX

DT 08-NOV-2000 (first entry)

XX

DE Peptide which binds to transcription factor E2F-1 DNA binding domain.

XX

KW DNA binding; transcription factor; E2F; E2F-1; cell cycle; DP-1; activation; transcription; apoptosis; proliferative disorder; psoriasis; restenosis.

KW

XX

OS Synthetic.

XX

PN WO200044771-A1.

XX

PD 03-AUG-2000.

XX

PF 26-JAN-2000; 2000WO-GB00227.

XX

PR 26-JAN-1999; 99GB-0001710.

XX

PA (PROL-) PROLIFIX LTD.

XX

PI Mueller R, Kontermann RE, Montigiani S;

XX

DR WPI; 2000-532806/48.

XX

PT Peptides binding to the DNA binding domain of transcription factor E2F and inhibiting cell cycle progression, useful for the treatment of cancer

PT

XX

PS Claim 6; Page 2; 42pp; English.

XX

CC Peptides which bind to the DNA binding domain of transcription factor E2F and inhibit cell cycle progression may be useful as research agents to investigate the interaction between E2F and DP-1, or the activation of transcription by E2F-1/DP-1 heterodimers. They may also be used for inducing apoptosis and/or cell cycle arrest in a cell, particularly for treatment of cancer or other proliferative disorders such as psoriasis and restenosis.

XX

SQ Sequence 6 AA;

Query Match 89.5%; Score 34; DB 21; Length 6;

Best Local Similarity 66.7%; Pred. No. 9.3e+05;

Matches 4; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

OY 1 WXXWHF 6

Db 1 WVRWHF 6

RESULT 4

AAB01499

ID AAB01499 standard; peptide; 6 AA.

XX

AC AAB01499;

XX

DT 08-NOV-2000 (first entry)

XX

DE Peptide which binds to transcription factor E2F-1 DNA binding domain.

XX

KW DNA binding; transcription factor; E2F; E2F-1; cell cycle; DP-1; activation; transcription; apoptosis; proliferative disorder; psoriasis; restenosis.

KW

XX OS Synthetic.  
XX FH Key Location/Qualifiers  
FT FT Misc-difference 2 /note= "Any amino acid"  
FT FT Misc-difference 3 /note= "Any amino acid"  
XX XX  
PN WO2000044771-A1.  
XX XX  
PD 03-AUG-2000.  
XX XX  
PF 26-JAN-2000; 2000WO-GB00227.  
XX XX  
PR 26-JAN-1999; 99GB-0001710.  
XX PA (PROL-) PROLIFIX LTD.  
XX PI Mueller R, Kontermann RE, Montigiani S;  
XX XX  
DR WPI; 2000-532806/48.  
XX XX  
PT Peptides binding to the DNA binding domain of transcription factor E2F  
PT and inhibiting cell cycle progression, useful for the treatment of  
PT cancer  
XX XX  
PS Claim 4; Page 9; 42pp; English.  
XX XX  
CC Peptides which bind to the DNA binding domain of transcription  
CC factor E2F and inhibit cell cycle progression may be useful as  
CC research agents to investigate the interaction between E2F and DP-1,  
CC or the activation of transcription by E2F-1/DP-1 heterodimers. They  
CC may also be used for inducing apoptosis and/or cell cycle arrest in  
CC a cell, particularly for treatment of cancer or other proliferative  
CC disorders such as psoriasis and restenosis.  
XX XX  
SQ Sequence 6 AA;

Query Match 89.5%; Score 34; DB 21; Length 6;  
Best Local Similarity 100.0%; Pred. No. 9.3e+05;  
Matches 6; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 WXXWHF 6  
| | | | |  
Db 1 WXXWHF 6

RESULT 5  
ABR45593  
ID ABR45593 standard; Peptide; 6 AA.  
XX XX  
AC ABR45593;  
XX XX  
DT 10-JUN-2003 (first entry)  
XX XX  
DE Staphylococcus aureus CHIPS-related peptide #783.  
XX XX  
KW CHIPS; Chemotaxis Inhibitory Protein; C5a-receptor; C5aR;  
KW formulated peptide receptor; FPR; neutrophil; monocyte; endothelial cell;  
KW inflammation; cardiovascular disease; central nervous system disease;  
KW gastrointestinal disease; skin disease; genitourinary disease;  
KW joint disease; respiratory disease; HIV infection; antiinflammatory;  
KW cardiant; cerebroprotective; neuroprotective; nootropic; dermatological;  
KW gynecological; immunosuppressive; anti-HIV.  
XX XX  
OS Staphylococcus aureus.  
OS Synthetic.  
XX XX  
PN WO2003006048-A1.  
XX XX  
PD 23-JAN-2003.

PF 11-JUL-2001; 2001WO-EP08004.  
XX XX  
PR 11-JUL-2001; 2001WO-EP08004.  
XX XX  
PA (JARI-) JARI PHARM BV.  
XX XX  
PI Van Kessel CPM, Gosselaar-de Haas CJC, Kruijtzer JAW;  
PI Van Strijp JAG;  
XX XX  
DR WPI; 2003-247783/25.  
XX XX  
PT Combination of peptides derived from chemotaxis inhibiting protein from  
PT Staphylococcus aureus (CHIPS) having CHIPS activity, useful in  
PT prophylaxis and treatment of inflammation, cardiovascular, skin and  
PT kidney diseases  
XX XX  
PS Disclosure; Page 13; 89pp; English.  
XX XX  
CC The present invention relates to peptides (ABR44811-ABR47162 and  
CC ABR47164-ABR47385) derived from the Chemotaxis Inhibitory Protein (CHIPS)  
CC from Staphylococcus aureus. The peptide fragments are useful in the  
CC prophylaxis or treatment of diseases or disorders involving the  
CC C5a-receptor (C5aR) and/or formulated peptide receptor (FPR) or  
CC neutrophils, monocytes and endothelial cells or involving acute or  
CC chronic inflammation reactions. The diseases or disorders include  
CC cardiovascular diseases, disease of the central nervous system,  
CC gastrointestinal diseases, skin diseases, genitourinary diseases, joint  
CC diseases, respiratory diseases and HIV infection.  
XX XX  
SQ Sequence 6 AA;

Query Match 76.3%; Score 29; DB 24; Length 6;  
Best Local Similarity 50.0%; Pred. No. 9.3e+05;  
Matches 3; Conservative 1; Mismatches 2; Indels 0; Gaps 0;

QY 1 WXXWHF 6  
| | | | |  
Db 1 WSWFYF 6

RESULT 6  
ABR45594  
ID ABR45594 standard; Peptide; 6 AA.  
XX XX  
AC ABR45594;  
XX XX  
DT 10-JUN-2003 (first entry)  
XX XX  
DE Staphylococcus aureus CHIPS-related peptide #784.  
XX XX  
KW CHIPS; Chemotaxis Inhibitory Protein; C5a-receptor; C5aR;  
KW formulated peptide receptor; FPR; neutrophil; monocyte; endothelial cell;  
KW inflammation; cardiovascular disease; central nervous system disease;  
KW gastrointestinal disease; skin disease; genitourinary disease;  
KW joint disease; respiratory disease; HIV infection; antiinflammatory;  
KW cardiant; cerebroprotective; neuroprotective; nootropic; dermatological;  
KW gynecological; immunosuppressive; anti-HIV.  
XX XX  
OS Staphylococcus aureus.  
OS Synthetic.  
XX XX  
PN WO2003006048-A1.  
XX XX  
PD 23-JAN-2003.  
XX XX  
PF 11-JUL-2001; 2001WO-EP08004.  
XX XX  
PR 11-JUL-2001; 2001WO-EP08004.  
XX XX  
PA (JARI-) JARI PHARM BV.  
XX XX  
PI Van Kessel CPM, Gosselaar-de Haas CJC, Kruijtzer JAW;  
PI Van Strijp JAG;

XX WPI; 2003-247783/25.

XX Combination of peptides derived from chemotaxis inhibiting protein from

PT Staphylococcus aureus (CHIPS) having CHIPS activity, useful in

PT prophylaxis and treatment of inflammation, cardiovascular, skin and

PT kidney diseases

XX Disclosure; Page 13; 89pp; English.

XX The present invention relates to peptides (ABR44811-ABR47162 and

CC ABR47164-ABR47385) derived from the Chemotaxis Inhibitory Protein (CHIPS)

CC from Staphylococcus aureus. The peptide fragments are useful in the

CC prophylaxis or treatment of diseases or disorders involving the

CC C5a-receptor (C5aR) and/or formylated peptide receptor (FPR) or

CC neutrophils, monocytes and endothelial cells or involving acute or

CC chronic inflammation reactions. The diseases or disorders include

CC cardiovascular diseases, disease of the central nervous system,

CC gastrointestinal diseases, skin diseases, genitourinary diseases, joint

CC diseases, respiratory diseases and HIV infection.

XX Sequence 6 AA;

Query Match 76.3%; Score 29; DB 24; Length 6;

Best Local Similarity 50.0%; Pred. No. 9.3e+05;

Matches 3; Conservative 1; Mismatches 2; Indels 0; Gaps 0;

OY 1 WXXWHP 6

Db 1 WTFWYF 6

RESULT 7

AAM00214

ID AAM00214 standard; Peptide; 14 AA.

XX AC AAM00214;

XX DT 01-OCT-2001 (first entry)

XX DE Human angiotensin fragment SEQ ID NO: 754.

XX KW Human; single nucleotide polymorphism; SNP; paternity test;

XX KW forensic test; aberrant protein expression.

XX OS Homo sapiens.

XX PN WO200151670-A2.

XX PD 19-JUL-2001.

XX PF 05-JAN-2001; 2001WO-US00322.

XX PR 07-JAN-2000; 2000US-0174962.

XX PA (CURA-) CURAGEN CORP.

XX PI Shimkets RA, Leach MD;

XX WPI; 2001-451871/48.

DR N-PSDB; AAH89323.

XX Isolated human polynucleotides containing single nucleotide

PT polymorphisms, useful for the treatment and diagnosis of e.g. cancer,

PT infection and diabetes

XX Disclosure; Page 321; 475pp; English.

XX The present invention relates to human nucleic acids containing single

CC nucleotide polymorphisms (SNPs). These can be used in forensic and

CC paternity tests, and to aid in the treatment of diseases associated with

CC aberrant protein expression, including cancer, amyloidosis, diabetes,

CC Alzheimer's disease, Down's syndrome, oedema, lupus (SLE), vasculitis,

CC glomerulonephritis, haemolytic anaemia, thrombocytopaenia, arthritis,

CC meningitis, muscular disorders, dementia, neurological diseases, tubercous

CC sclerosis, male infertility, hypercalcaemia, blood pressure disorders,

CC osteoporosis, pathogenic infections, hypercholesterolaemia, obesity or

CC autoimmunity. The present sequence is a peptide encoded by a

CC polymorphism-containing oligonucleotide fragment of the invention.

XX Sequence 14 AA;

Query Match 76.3%; Score 29; DB 22; Length 14;

Best Local Similarity 60.0%; Pred. No. 1.1e+02;

Matches 3; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

OY 1 WXXWH 5

Db 7 WYTW 11

RESULT 8

AAW28912

ID AAW28912 standard; peptide; 6 AA.

XX AC AAW28912;

XX DT 20-JAN-1998 (first entry)

XX DE Opioid peptide.

XX KW enkephalin; mu-opioid receptor ligand; agonist; antagonist.

XX OS Synthetic.

XX FH Key Location/Qualifiers

FT Modified-site 1

FT Modified-site /note= "N-acetyl-Arg"

FT Modified-site 6

FT /note= "the C-terminal is in amide form"

XX PN US5641861-A.

XX PD 24-JUN-1997.

XX PF 07-JUN-1995; 95US-0487006.

XX PR 07-JUN-1995; 95US-0487006.

XX (TORR-) TORREY PINES INST MOLECULAR STUDIES.

XX PA Dooley CT, Houghten RA;

XX WPI; 1997-340994/31.

XX New opioid peptides which bind mu receptors specifically - have

PT agonist or antagonist activity and are used for study and

PT localisation of mu receptors and to treat peripheral side effects of

PT morphine etc.

XX Disclosure; Column 8; 92pp; English.

XX The patent discloses the following new peptides, which are opioids which

CC bind specifically to the mu receptor: Ac-Phe-Arg-Trp-Trp-Tyr-Xaa-NH2 (1);

CC Ac-Arg-Trp-Ile-Gly-Trp-Xaa-NH2 (2); Trp-Trp-Pro-Lys-His-Xaa-NH2 (3);

CC Trp-Trp-Pro-Xaa-NH2 (4); Tyr-Pro-Phe-Gly-Phe-Xaa-NH2 (5);

CC D-Ile-D-Met-D-Ser-D-Trp-D-Trp-(Gly)n-Xaa2-NH2 (6);

CC D-Ile-D-Met-D-Thr-D-Trp-Gly-Xaa2-NH2 (7); Tyr-Al-B2-C3-NH2 (214);

CC Pm and red ((Me)x(H)y-Tyr-(NMe)z-Xaa3)z-NH2 (221); and

CC Trp-Trp-Pro-D4-(His)z-(Xaa)z-NH2 (222); where Xaa = any natural amino

CC acid; Xaa1 = Lys or Arg; n and z = 0 or 1; Xaa2 = Gly or the D form of

CC any naturally occurring amino acid; Al = D-norvaline or D-norleucine;

CC B2 = Gly, Phe or Trp; C3 = Trp or naphthylalanine; x and y = 0-2, but

CC not over 2 in total; Xaa3 = Phe, DPhe or benzylamino; D4 = Lys or Arg;

CC Pm and red indicate permethylation and reduction of all CO in peptide

CC links to methylene. These new compounds are useful: (i) for in vitro



CC assay and study of opiate receptor subtypes, particularly mu receptors  
CC in the brain; (ii) for in vivo localisation of receptor subtypes; and  
CC (iii) therapeutically to block the peripheral effects (e.g. constipation  
CC and pruritus) of centrally acting pain killers such as morphine.  
CC They are very selective for the mu opioid receptor, over binding to the  
CC delta and kappa receptor subtypes.  
CC The present sequence is a specific example of a peptide (2).

XX Sequence 6 AA;  
Query Match 73.7%; Score 28; DB 18; Length 6;  
Best Local Similarity 60.0%; Pred. No. 9.3e+05;  
Matches 3; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

QY 1 WXXWH 5  
Db 2 WIGWH 6  
RESULT 9  
AAR93770  
ID AAR93770 standard; Protein; 6 AA.  
XX  
AC AAR93770;  
XX  
DT 23-SEP-1997 (first entry)  
XX  
DE New peptide which acts as mu-opioid receptor ligand.  
XX  
KW mu-receptor; opioid; opiate; agonist; antagonist; diagnosis;  
KW analgesic.  
XX  
OS Synthetic.  
XX  
FH Key Location/Qualifiers  
FT Modified-site 1 /note= "N-acetyl-Arg"  
FT Misc-difference 6 /note= "this residue is in C-terminal amide form"

XX WO9640208-A1.  
XX  
PD 19-DEC-1996.  
XX  
PF 06-JUN-1996; 96WO-US09321.  
XX  
PR 07-JUN-1995; 95US-0476438.  
XX  
PA (TORR-) TORREY PINES INST MOLECULAR STUDIES.  
XX  
PI Dooley CT, Houghten RA;  
DR WPI; 1997-051895/05.  
XX  
PT New mu opioid receptor binding ligand peptide(s) - useful for  
PT in-vitro and in-vivo diagnosis, as analgesics, and for blocking  
PT peripheral effects of centrally acting drugs, e.g. morphine  
XX  
PS Disclosure; Page 19; 57pp; English.  
XX  
CC The patent discloses eight new groups of opioid peptides which bind  
CC to the mu-receptor to act as agonists or antagonists. The peptides  
CC can be used for in-vitro assays to study opiate receptor subtypes  
CC (especially the mu type) in brain or other tissue samples; and for  
CC in-vivo diagnosis to localise opioid subtypes. The peptides are also  
CC useful as drugs to treat pathologies associated with other compounds  
CC which interact with the opioid receptor system. Therefore they can be  
CC used in medicaments for treating pathologies associated with the mu  
CC receptor and as analgesics. They can be used therapeutically to block  
CC the peripheral effects of centrally acting pain killers, e.g. to  
CC prevent side effects such as constipation and pruritis associated  
CC with morphine. The present sequence represents a specific example  
CC of one of the new groups of peptides, of formula

CC Ac-Arg-Trp-Ile-Gly-Trp-Xaa-NH2 where Xaa = a naturally occurring  
CC amino acid.  
XX  
SQ Sequence 6 AA;  
Query Match 73.7%; Score 28; DB 18; Length 6;  
Best Local Similarity 60.0%; Pred. No. 9.3e+05;  
Matches 3; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

QY 1 WXXWH 5  
Db 2 WIGWH 6  
RESULT 10  
AAY23019  
ID AAY23019 standard; peptide; 6 AA.  
XX  
AC AAY23019;  
XX  
DT 23-AUG-1999 (first entry)  
XX  
DE Opioid peptide which inhibits binding of enkephalin.  
XX  
KW Opioid peptide; ligand binding; opioid receptor;  
KW micro-selective opioid peptide; enkephalin; opioid receptor system;  
KW blocking; peripheral effect; centrally acting pain killer; morphine.  
XX  
OS Synthetic.  
XX  
FH Key Location/Qualifiers  
FT Modified-site 1 /note= "acetylated"  
FT Modified-site 6 /note= "amidated"

XX US5919897-A.  
XX  
PD 06-JUL-1999.  
XX  
PF 07-JUN-1995; 95US-0488659.  
XX  
PR 07-JUN-1995; 95US-0488659.  
XX  
PA (TORR-) TORREY PINES INST MOLECULAR STUDIES.  
XX  
PI Dooley CT, Houghten RA;  
DR WPI; 1999-394647/33.  
XX  
PT New opioid peptides useful for blocking the peripheral effects of  
PT centrally acting pain killers such as morphine  
XX  
PS Example 1; Column 8; 92pp; English.  
XX  
CC The specification describes opioid peptides, in which each of the  
CC N atoms in the peptide backbone between respective amino acids is  
CC modified by permethylation, perallylation, perethylation, perbenzylation.  
CC and pernaphthylation. The peptides inhibit ligand binding to an opioid  
CC receptor. Specifically, the peptides inhibit the micro-selective  
CC opioid peptide enkephalin. The peptides can be used in vivo  
CC diagnostically to localise opioid receptor subtypes. They can be used  
CC to treat pathologies associated with other compounds which interact with  
CC the opioid receptor system. The peptides are especially useful for  
CC blocking the peripheral effects of centrally acting pain killers such  
CC as morphine. AAY23005-Y23024 represent opioid peptides of the invention,  
CC and are derived from the general sequence given in AAY23004.

XX  
SQ Sequence 6 AA;  
Query Match 73.7%; Score 28; DB 20; Length 6;  
Best Local Similarity 60.0%; Pred. No. 9.3e+05;  
Matches 3; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

QY 1 WXXWH 5  
Db 2 WIGWH 6

RESULT 11  
AAB01509  
ID AAB01509 standard; peptide; 6 AA.  
XX AC AAB01509;  
XX DT 08-NOV-2000 (first entry)  
XX DE Peptide which binds to transcription factor E2F-1 DNA binding domain.  
XX KW DNA binding; transcription factor; E2F; E2F-1; cell cycle; DP-1;  
KW activation; transcription; apoptosis; proliferative disorder;  
KW psoriasis; restenosis.  
XX OS Synthetic.  
XX PN WO200044771-A1.  
XX PD 03-AUG-2000.  
XX PF 26-JAN-2000; 2000WO-GB00227.  
XX PR 26-JAN-1999; 99GB-0001710.  
XX PA (PROL-). PROLIFIX LTD.  
XX PI Mueller R, Kontermann RE, Montigiani S;  
XX DR WPI; 2000-532806/48.  
XX PT Peptides binding to the DNA binding domain of transcription factor E2F  
PT and inhibiting cell cycle progression, useful for the treatment of  
PT cancer  
XX PS Example; Page 26; 42pp; English.  
XX CC Peptides which bind to the DNA binding domain of transcription  
CC factor E2F and inhibit cell cycle progression may be useful as  
CC research agents to investigate the interaction between E2F and DP-1,  
CC or the activation of transcription by E2F-1/DP-1 heterodimers. They  
CC may also be used for inducing apoptosis and/or cell cycle arrest in  
CC a cell, particularly for treatment of cancer or other proliferative  
CC disorders such as psoriasis and restenosis.  
XX SQ Sequence 6 AA;

Query Match 73.7%; Score 28; DB 21; Length 6;  
Best Local Similarity 60.0%; Pred. No. 9.3e+05;  
Matches 3; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

QY 1 WXXWH 5  
Db 1 WVRWH 5

RESULT 12  
ABR45591  
ID ABR45591 standard; Peptide; 6 AA.  
XX AC ABR45591;  
XX DT 10-JUN-2003 (first entry)  
XX DE Staphylococcus aureus CHIPS-related peptide #781.  
XX KW CHIPS; Chemotaxis Inhibitory Protein; C5a-receptor; C5aR;  
KW formylated peptide receptor; FPR; neutrophil; monocyte; endothelial cell;

KW inflammation; cardiovascular disease; central nervous system disease;  
KW gastrointestinal disease; skin disease; genitourinary disease;  
KW joint disease; respiratory disease; HIV infection; antiinflammatory;  
KW cardiant; cerebroprotective; neuroprotective; nootropic; dermatological;  
KW gynecological; immunosuppressive; anti-HIV.  
XX Staphylococcus aureus.  
OS Synthetic.  
OS XX WO2003006048-A1.  
XX PD 23-JAN-2003.  
XX PF 11-JUL-2001; 2001WO-EP08004.  
XX PR 11-JUL-2001; 2001WO-EP08004.  
XX PA (JARI-) JARI PHARM BV.  
XX Van Kessel CPM, Gosselaar-de Haas CJC, Kruijtzer JAW;  
PI Van Strijp JAG;  
PI XX WPI; 2003-247783/25.  
DR XX Combination of peptides derived from chemotaxis inhibiting protein from  
XX Staphylococcus aureus (CHIPS) having CHIPS activity, useful in  
PT prophylaxis and treatment of inflammation, cardiovascular, skin and  
PT kidney diseases  
XX PS Disclosure; Page 13; 89pp; English.  
XX CC The present invention relates to peptides (ABR44811-ABR47162 and  
CC ABR47164-ABR47385) derived from the Chemotaxis Inhibitory Protein (CHIPS)  
CC from Staphylococcus aureus. The peptide fragments are useful in the  
CC prophylaxis or treatment of diseases or disorders involving the  
CC C5a-receptor (C5aR) and/or formylated peptide receptor (FPR) or  
CC neutrophils, monocytes and endothelial cells or involving acute or  
CC chronic inflammation reactions. The diseases or disorders include  
CC cardiovascular diseases, disease of the central nervous system,  
CC gastrointestinal diseases, skin diseases, genitourinary diseases, joint  
CC diseases, respiratory diseases and HIV infection.  
XX SQ Sequence 6 AA;

Query Match 73.7%; Score 28; DB 24; Length 6;  
Best Local Similarity 50.0%; Pred. No. 9.3e+05;  
Matches 3; Conservative 1; Mismatches 2; Indels 0; Gaps 0;

QY 1 WXXWHF 6  
Db 1 WFFWYF 6

RESULT 13  
ABR45592  
ID ABR45592 standard; Peptide; 6 AA.  
XX AC ABR45592;  
XX DT 10-JUN-2003 (first entry)  
XX DE Staphylococcus aureus CHIPS-related peptide #782.  
XX KW CHIPS; Chemotaxis Inhibitory Protein; C5a-receptor; C5aR;  
KW formylated peptide receptor; FPR; neutrophil; monocyte; endothelial cell;  
KW inflammation; cardiovascular disease; central nervous system disease;  
KW gastrointestinal disease; skin disease; genitourinary disease;  
KW joint disease; respiratory disease; HIV infection; antiinflammatory;  
KW cardiant; cerebroprotective; neuroprotective; nootropic; dermatological;  
KW gynecological; immunosuppressive; anti-HIV.  
XX Staphylococcus aureus.  
OS Synthetic.

XX WO2003006048-A1.  
PN  
XX  
PD 23-JAN-2003.  
XX  
PF 11-JUL-2001; 2001WO-EP08004.  
XX  
PR 11-JUL-2001; 2001WO-EP08004.  
XX  
PA (JARI-) JARI PHARM BV.  
XX  
PI Van Kessel CPM, Gosselaar-de Haas CJC, Kruijtzer JAW;  
PI Van Strijp JAG;  
XX  
DR WPI; 2003-247783/25.  
XX  
PT Combination of peptides derived from chemotaxis inhibiting protein from  
PT Staphylococcus aureus (CHIPS) having CHIPS activity, useful in  
PT prophylaxis and treatment of inflammation, cardiovascular, skin and  
PT kidney diseases  
XX  
PS Disclosure; Page 13; 89pp; English.  
XX  
CC The present invention relates to peptides (ABR44811-ABR47162 and  
CC ABR47164-ABR47385) derived from the Chemotaxis Inhibitory Protein (CHIPS)  
CC from Staphylococcus aureus. The peptide fragments are useful in the  
CC prophylaxis or treatment of diseases or disorders involving the  
CC C5a-receptor (C5aR) and/or formylated peptide receptor (FPR) or  
CC neutrophils, monocytes and endothelial cells or involving acute or  
CC chronic inflammation reactions. The diseases or disorders include  
CC cardiovascular diseases, disease of the central nervous system,  
CC gastrointestinal diseases, skin diseases, genitourinary diseases, joint  
CC diseases, respiratory diseases and HIV infection.  
XX  
SQ Sequence 6 AA;  
  
Query Match 73.7%; Score 28; DB 24; Length 6;  
Best Local Similarity 50.0%; Pred. No. 9.3e+05;  
Matches 3; Conservative 1; Mismatches 2; Indels 0; Gaps 0;  
  
QY 1 WXXWHF 6  
Db 1 WIFWYF 6  
  
RESULT 14  
AAY01258  
ID AAY01258 standard; peptide; 7 AA.  
XX  
AC AAY01258;  
XX  
DT 01-JUN-1999 (first entry)  
XX  
DE US5851813 peptide sequence number 45.  
XX  
KW Antigenic composition; primate; lentivirus; nef gene; vaccine;  
KW infection; AIDS; SIVmac239; deletion; mutant.  
XX  
OS Simian immunodeficiency virus.  
OS Synthetic.  
XX  
PN US5851813-A.  
XX  
PD 22-DEC-1998.  
XX  
PF 27-JAN-1994; 94US-0188583.  
XX  
PR 27-JAN-1994; 94US-0188583.  
PR 12-JUL-1990; 90US-0551945.  
PR 09-JUL-1991; 91US-0727494.  
XX  
PA (HARD ) HARVARD COLLEGE.  
XX

PI Desrosiers RC;  
XX  
DR WPI; 1999-080408/07.  
DR N-PSDB; AAX27657.  
XX  
PT Lentivirus antigenic compositions - containing lentivirus with nef  
PT gene deletion  
XX  
PS Disclosure; Fig 5A-B; 93pp; English.  
XX  
CC The invention relates to an antigenic composition comprising an isolated  
CC primate lentivirus whose genome contains an engineered non-revertible  
CC null mutation in the nef gene, or an infectious DNA clone in a carrier.  
CC The antigenic composition is used in vaccines against infection by the  
CC lentivirus, e.g. AIDS.  
XX  
SQ Sequence 7 AA;  
  
Query Match 73.7%; Score 28; DB 20; Length 7;  
Best Local Similarity 60.0%; Pred. No. 9.3e+05;  
Matches 3; Conservative 0; Mismatches 2; Indels 0; Gaps 0;  
  
QY 1 WXXWH 5  
Db 1 WEYWH 5  
  
RESULT 15  
AAB49729  
ID AAB49729 standard; peptide; 7 AA.  
XX  
AC AAB49729;  
XX  
DT 10-APR-2001 (first entry)  
XX  
DE Peptide SEQ ID 40 which binds to the TADG5 protein.  
XX  
KW TADG5; human; zinc finger; SH3 domain; cell signalling;  
KW cell cycle control.  
XX  
OS Unidentified.  
XX  
PN WO200102432-A1.  
XX  
PD 11-JAN-2001.  
XX  
PF 30-JUN-2000; 2000WO-US18304.  
XX  
PR 01-JUL-1999; 99US-0346510.  
XX  
PA (UYAR-) UNIV ARKANSAS.  
XX  
PI O'Brien TJ, Wang Y;  
XX  
DR WPI; 2001-123102/13.  
XX  
PT Novel SH3 domain-containing TADG5 protein useful for regulating gene  
PT replication, as a nutrition supplement, and as a marker for human  
PT tissue, or in cell cycle control -  
XX  
PS Example 6; Page 36; 85pp; English.  
XX  
CC This invention relates to an SH3 domain-containing protein termed TADG5,  
CC and its variants. The invention includes amino acid and polynucleotide  
CC sequences for TADG5, and oligonucleotides which bind to either the basic  
CC amino acid region and/or the zinc finger motif of the TADG5 protein. The  
CC basic amino acid region or zinc finger motif of TADG5 is useful for  
CC regulating the expression of the TADG5 gene in a cell. The TADG5 protein  
CC is useful as a source of amino acids, as a nutrition supplement, and as a  
CC marker for human tissue, or in cell cycle control. TADG5 protein or  
CC peptides generated from the protein sequence are useful as antigens for  
CC the production of polyclonal and monoclonal antibodies. DNA encoding  
CC TADG5 is useful as an antisense vehicle for cell cycle control by

CC shutting down signalling or cell division. The present sequence  
CC represents a peptide identified from a phage display peptide library  
CC through biopanning with the TADG5 protein.

XX

SQ Sequence 7 AA;

Query Match 73.7%; Score 28; DB 22; Length 7;  
Best Local Similarity 60.0%; Pred. No. 9.3e+05;  
Matches 3; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

QY 1 WXXWH 5

| | |

Db 3 WMDWH 7

Search completed: December 3, 2003, 11:51:16

Job time : 34.6667 secs

GenCore version 5.1.6  
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OM protein - protein search, using sw model  
Run on: December 3, 2003, 11:48:35 ; Search time 11 Seconds  
(without alignments)  
52.456 Million cell updates/sec

Title: US-09-912-414-11  
Perfect score: 38  
Sequence: 1 WXXWHF 6

Scoring table: BLOSUM62  
Gapop 10.0 , Gapext 0.5

Searched: 283308 seqs, 96168682 residues

Total number of hits satisfying chosen parameters: 2520

Minimum DB seq length: 0  
Maximum DB seq length: 15

Post-processing: Minimum Match 0%  
Maximum Match 100%  
Listing first 45 summaries

Database : PIR\_76:\*  
1: pir1:\*  
2: pir2:\*  
3: pir3:\*  
4: pir4:\*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match %	Length	DB	ID	Description
1	23	60.5	9	2	S07241	litorin - Rohde's
2	22	57.9	9	2	S07205	litorin 2-Glu - Au
3	22	57.9	9	2	S07204	litorin I - Austro
4	22	57.9	10	2	F49033	T-cell receptor ga
5	22	57.9	13	2	A60409	bombesin-like pept
6	21	55.3	9	2	A43848	cell surface adhes
7	21	55.3	12	2	PH1308	Ig heavy chain DJ
8	20	52.6	12	2	PH1324	Ig heavy chain DJ
9	20	52.6	13	2	S61798	T-cell-specific tr
10	20	52.6	14	2	PH1322	Ig heavy chain DJ
11	18	47.4	9	2	D57444	neuropeptide Grb-A
12	17	44.7	12	2	A29169	phospholipase A2 (
13	17	44.7	15	2	PA0099	phenotypic variati
14	16	42.1	8	2	T13818	cytochrome oxidase
15	16	42.1	10	2	PQ0177	neuromedin C - lau
16	16	42.1	10	2	A60647	neuromedin C - bov
17	16	42.1	10	2	T13976	cytochrome-c oxida
18	16	42.1	10	2	T17057	cytochrome-c oxida
19	16	42.1	10	2	T12303	cytochrome-c oxida
20	16	42.1	10	2	T14019	cytochrome-c oxida
21	16	42.1	10	2	T17060	cytochrome-c oxida
22	16	42.1	10	2	T14043	cytochrome-c oxida
23	16	42.1	10	2	T14054	cytochrome-c oxida
24	16	42.1	10	2	T17066	cytochrome-c oxida
25	16	42.1	10	2	T17069	cytochrome-c oxida
26	16	42.1	10	2	T12308	cytochrome-c oxida
27	16	42.1	10	2	T17072	cytochrome-c oxida
28	16	42.1	10	2	T12312	cytochrome-c oxida
29	16	42.1	10	2	T12316	cytochrome-c oxida

30	16	42.1	10	2	T12321	cytochrome-c oxida
31	16	42.1	10	2	T14219	cytochrome-c oxida
32	16	42.1	14	1	BSTD	bombesin - fire-be
33	16	42.1	14	2	PT0077	proteochoondoitin c
34	15	39.5	9	2	S56004	glucan 1,3-beta-gl
35	15	39.5	12	2	S25039	Ig heavy chain V r
36	15	39.5	13	2	S23372	T-cell receptor al
37	15	39.5	13	2	B25448	Ig kappa-1 chain,
38	15	39.5	13	2	B26406	Ig kappa chain J r
39	15	39.5	13	2	A47630	Ig kappa chain J r
40	15	39.5	15	2	S24159	leukocyte elastase
41	14	36.8	7	2	S21230	dermorphin (Trp-4,
42	14	36.8	10	2	A58365	neuropeptide PPRFa
43	14	36.8	10	2	T17054	cytochrome-c oxida
44	14	36.8	10	2	T17063	cytochrome-c oxida
45	14	36.8	10	2	T12325	cytochrome-c oxida

ALIGNMENTS

RESULT 1

S07241  
litorin - Rohde's leaf frog  
C;Species: Phyllomedusa rohdei (Rohde's leaf frog)  
C;Date: 12-Feb-1993 #sequence\_revision 12-Mar-1993 #text\_change 18-Aug-2000  
C;Accession: S07241  
R;Barra, D.; Falconieri Erspamer, G.; Simmaco, M.; Bossa, F.; Melchiorri, P.; Erspamer  
FEBS Lett. 182, 53-56, 1985  
A;Title: Rohdei-litorin: a new peptide from the skin of Phyllomedusa rohdei.  
A;Reference number: S07241; MUID:85127560; PMID:3838283  
A;Accession: S07241  
A;Molecule type: protein  
A;Residues: 1-9 <BAR>  
C;Superfamily: gastrin-releasing peptide  
C;Keywords: amidated carboxyl end; blocked amino end; neuropeptide; pyroglutamic acid  
F;1/Modified site: pyrrolidone carboxylic acid (Gln) #status experimental  
F;9/Modified site: amidated carboxyl end (Met) #status experimental

Query Match 60.5%; Score 23; DB 2; Length 9;  
Best Local Similarity 50.0%; Pred. No. 2.8e+05;  
Matches 3; Conservative 0; Mismatches 3; Indels 0; Gaps 0;

QY 1 WXXWHF 6

Db 3 WATGHF 8

RESULT 2

S07205  
litorin 2-Glu - Australian tree frog (Litoria aurea)  
C;Species: Litoria aurea  
C;Date: 12-Feb-1993 #sequence\_revision 12-Mar-1993 #text\_change 18-Aug-2000  
C;Accession: S07205  
R;Anastasi, A.; Montecucchi, P.; Angelucci, F.; Erspamer, V.; Endean, R.  
Experientia 33, 1289, 1977  
A;Title: Glu(OMe)(2)-litorin, the second bombesin-like peptide occurring in methanol e:  
A;Reference number: S07205; MUID:78003546; PMID:908397  
A;Accession: S07205  
A;Molecule type: protein  
A;Residues: 1-9 <ANA>  
C;Superfamily: gastrin-releasing peptide  
C;Keywords: amidated carboxyl end; neuropeptide; pyroglutamic acid  
F;1/Modified site: pyrrolidone carboxylic acid (Gln) #status experimental  
F;9/Modified site: amidated carboxyl end (Met) #status experimental

Query Match 57.9%; Score 22; DB 2; Length 9;

Best Local Similarity 50.0%; Pred. No. 2.8e+05;

Matches 3; Conservative 0; Mismatches 3; Indels 0; Gaps 0;

QY 1 WXXWHF 6

Db 3 WATGHF 8



```
RESULT 3
S07204
litorin I - Australian tree frog (Litoria aurea)
C;Species: Litoria aurea
C;Date: 12-Feb-1993 #sequence_revision 12-Mar-1993 #text_change 18-Aug-2000
C;Accession: S07204
R;Anastasi, A.; Erspamer, V.; Endean, R.
Experientia 31, 510-511, 1975
A;Title: Aminoacid composition and sequence of litorin, a bombesin-like nonapeptide from
A;Reference number: S07204; MUID:75187011; PMID:1140241
A;Accession: S07204
A;Molecule type: protein
A;Residues: 1-9 <ANA>
C;Superfamily: gastrin-releasing peptide
C;Keywords: amidated carboxyl end; neuropeptide; pyroglutamic acid
F;1/Modified site: pyrrolidone carboxylic acid (Gln) #status experimental
F;9/Modified site: amidated carboxyl end (Met) #status experimental

Query Match      57.9%; Score 22; DB 2; Length 9;
Best Local Similarity 50.0%; Pred. No. 2.8e+05;
Matches 3; Conservative 0; Mismatches 3; Indels 0; Gaps 0;

QY      1 WXXWHF 6
      |  |  |
Db      3 WVGHF 8

RESULT 4
F49033
T-cell receptor gamma chain V-D-J region - human (fragment)
C;Species: Homo sapiens (man)
C;Date: 19-Dec-1993 #sequence_revision 17-Mar-2000 #text_change 17-Mar-2000
C;Accession: F49033
R;Morita, C.T.; Verma, S.; Aparicio, P.; Martinez, C.; Spits, H.; Brenner, M.B.
Eur. J. Immunol. 21, 2999-3007, 1991
A;Title: Functionally distinct subsets of human gamma/delta T cells.
A;Reference number: A49033; MUID:92083926; PMID:1684157
A;Accession: F49033
A;Status: preliminary
A;Molecule type: DNA
A;Residues: 1-10 <MOR>
A;Cross-references: GB:S72605; NID:G240700; PIDN:AAB20632.1; PID:G240701
A;Note: sequence extracted from NCBI backbone (NCBIN:72605, NCBIP:72606)
C;Keywords: T-cell receptor

Query Match      57.9%; Score 22; DB 2; Length 10;
Best Local Similarity 40.0%; Pred. No. 3.7e+02;
Matches 2; Conservative 1; Mismatches 2; Indels 0; Gaps 0;

QY      1 WXXWH 5
      |  |  |
Db      4 WERYW 8

RESULT 5
A60409
bombesin-like peptide L - frog (Pseudophryne guentheri)
C;Species: Pseudophryne guentheri
C;Date: 30-Jan-1993 #sequence_revision 30-Jan-1993 #text_change 18-Aug-2000
C;Accession: A60409
R;Simmaco, M.; Severini, C.; De Biase, D.; Barra, D.; Bossa, F.; Roberts, J.D.; Melchior
Peptides 11, 299-304, 1990
A;Title: Six novel tachykinin- and bombesin-related peptides from the skin of the Austr
A;Reference number: A60409; MUID:90287814; PMID:2356157
A;Accession: A60409
A;Molecule type: protein
A;Residues: 1-13 <SIM>
C;Superfamily: unassigned animal peptides
C;Keywords: amidated carboxyl end; hormone; neuropeptide; pyroglutamic acid
F;1/Modified site: pyrrolidone carboxylic acid (Gln) #status experimental
F;13/Modified site: amidated carboxyl end (Met) #status experimental
```

```
Query Match      57.9%; Score 22; DB 2; Length 13;
Best Local Similarity 50.0%; Pred. No. 4.6e+02;
Matches 3; Conservative 0; Mismatches 3; Indels 0; Gaps 0;

QY      1 WXXWHF 6
      |  |  |
Db      7 WVGHF 12

RESULT 6
A43848
cell surface adhesin for heparan sulfate, 66K - Staphylococcus aureus (fragment)
C;Species: Staphylococcus aureus
C;Date: 10-Mar-1993 #sequence_revision 18-Nov-1994 #text_change 24-Feb-1995
C;Accession: A43848
R;Liang, O.D.; Ascencio, F.; Fransson, L.A.; Wadstrom, T.
Infect. Immun. 60, 899-906, 1992
A;Title: Binding of heparan sulfate to Staphylococcus aureus.
A;Reference number: A43848; MUID:92176005; PMID:1541563
A;Accession: A43848
A;Status: preliminary
A;Molecule type: protein
A;Residues: 1-9 <LIA>
A;Note: sequence extracted from NCBI backbone (NCBIP:85442)

Query Match      55.3%; Score 21; DB 2; Length 9;
Best Local Similarity 50.0%; Pred. No. 2.8e+05;
Matches 2; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

QY      1 WXXW 4
      |  |  |
Db      2 WTGW 5

RESULT 7
PH1308
Ig heavy chain DJ region (clone C731-94) - human (fragment)
C;Species: Homo sapiens (man)
C;Date: 30-Sep-1993 #sequence_revision 30-Sep-1993 #text_change 07-May-1999
C;Accession: PH1308
R;Wasserman, R.; Galili, N.; Ito, Y.; Reichard, B.A.; Shane, S.; Rovera, G.
J. Exp. Med. 176, 1577-1581, 1992
A;Title: Predominance of fetal type DJH joining in young children with B precursor lym
A;Reference number: PH1302; MUID:93094761; PMID:1460419
A;Accession: PH1308
A;Molecule type: DNA
A;Residues: 1-12 <WAS>
C;Keywords: heterotetramer; immunoglobulin

Query Match      55.3%; Score 21; DB 2; Length 12;
Best Local Similarity 40.0%; Pred. No. 6.2e+02;
Matches 2; Conservative 1; Mismatches 2; Indels 0; Gaps 0;

QY      1 WXXWH 5
      |  |  |
Db      7 WQWN 11

RESULT 8
PH1324
Ig heavy chain DJ region (clone C510-100) - human (fragment)
C;Species: Homo sapiens (man)
C;Date: 30-Sep-1993 #sequence_revision 30-Sep-1993 #text_change 07-May-1999
C;Accession: PH1324
R;Wasserman, R.; Galili, N.; Ito, Y.; Reichard, B.A.; Shane, S.; Rovera, G.
J. Exp. Med. 176, 1577-1581, 1992
A;Title: Predominance of fetal type DJH joining in young children with B precursor lym
A;Reference number: PH1302; MUID:93094761; PMID:1460419
A;Accession: PH1324
A;Molecule type: DNA
A;Residues: 1-12 <WAS>
C;Keywords: heterotetramer; immunoglobulin
```

Query Match 52.6%; Score 20; DB 2; Length 12;  
Best Local Similarity 50.0%; Pred. No. 9e+02;  
Matches 2; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

QY 1 WXXW 4  
| |  
Db 5 WYW 8

RESULT 9  
S61798  
T-cell-specific transcription factor 1 splice form G - human (fragment)  
N;Alternate names: transcription factor TCF-1G  
C;Species: Homo sapiens (man)  
C;Date: 19-Mar-1997 #sequence\_revision 18-Jul-1997 #text\_change 24-Jul-1998  
C;Accession: S61798; S61880  
R;Mayer, K.; Wolff, E.; Clevers, H.; Ballhausen, W.G.  
Biochim. Biophys. Acta 1263, 169-172, 1995  
A;Title: The human high mobility group (HMG)-box transcription factor TCF-1: novel isoform  
A;Reference number: S61796; MUID:95367594; PMID:7640309  
A;Accession: S61798  
A;Molecule type: mRNA  
A;Residues: 1-13 <MAY>  
A;Cross-references: EMBL:Z47364  
A;Note: DNA was also sequenced  
C;Keywords: alternative splicing; DNA binding; transcription factor

Query Match 52.6%; Score 20; DB 2; Length 13;  
Best Local Similarity 50.0%; Pred. No. 9.7e+02;  
Matches 2; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

QY -1 WXXW 4  
| |  
Db 6 WDGW 9

RESULT 10  
PH1322  
Ig heavy chain DJ region (clone C344-99) - human (fragment)  
C;Species: Homo sapiens (man)  
C;Date: 30-Sep-1993 #sequence\_revision 30-Sep-1993 #text\_change 07-May-1999  
C;Accession: PH1322  
R;Wasserman, R.; Galili, N.; Ito, Y.; Reichard, B.A.; Shane, S.; Rovera, G.  
J. Exp. Med. 176, 1577-1581, 1992  
A;Title: Predominance of fetal type DJH joining in young children with B precursor lymphoma  
A;Reference number: PH1302; MUID:93094761; PMID:1460419  
A;Accession: PH1322  
A;Molecule type: DNA  
A;Residues: 1-14 <WAS>  
C;Keywords: heterotetramer; immunoglobulin

Query Match 52.6%; Score 20; DB 2; Length 14;  
Best Local Similarity 50.0%; Pred. No. 1e+03;  
Matches 2; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

QY 1 WXXW 4  
| |  
Db 6 WDW 9

RESULT 11  
D57444  
neuropeptide Grb-AST B4 - two-spotted cricket  
C;Species: Gryllus bimaculatus (two-spotted cricket)  
C;Date: 26-Jan-1996 #sequence\_revision 26-Jan-1996 #text\_change 26-Jan-1996  
C;Accession: D57444  
R;Lorenz, M.W.; Kellner, R.; Hoffmann, K.H.  
J. Biol. Chem. 270, 21103-21108, 1995  
A;Title: A family of neuropeptides that inhibit juvenile hormone biosynthesis in the cricket  
A;Reference number: A57444; MUID:95403341; PMID:7673141  
A;Accession: D57444  
A;Status: preliminary

A;Molecule type: protein  
A;Residues: 1-9 <LOR>

Query Match 47.4%; Score 18; DB 2; Length 9;  
Best Local Similarity 40.0%; Pred. No. 2.8e+05;  
Matches 2; Conservative 1; Mismatches 2; Indels 0; Gaps 0;

QY 1 WXXWH 5  
| |  
Db 2 WERFH 6

RESULT 12  
A29169  
phospholipase A2 (EC 3.1.1.4) precursor - sheep (fragment)  
C;Species: Ovis orientalis aries, Ovis ammon aries (domestic sheep)  
C;Date: 02-Jun-1988 #sequence\_revision 02-Jun-1988 #text\_change 31-Oct-1997  
C;Accession: A29169  
R;Dutilh, C.E.; Van Doren, P.J.; Verheul, F.E.A.M.; De Haas, G.H.  
Eur. J. Biochem. 53, 91-97, 1975  
A;Title: Isolation and properties of phospholipase A2 from ox and sheep pancreas.  
A;Reference number: A94661  
A;Accession: A29169  
A;Molecule type: protein  
A;Residues: 1-12 <DUT>  
C;Superfamily: phospholipase A2  
C;Keywords: carboxylic ester hydrolase; pyroglutamic acid  
F;1/Modified site: pyrrolidone carboxylic acid (Gln) #status experimental

Query Match 44.7%; Score 17; DB 2; Length 12;  
Best Local Similarity 66.7%; Pred. No. 2.8e+03;  
Matches 2; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

QY 4 WHF 6  
| |  
Db 10 WQF 12

RESULT 13  
PA0099  
phenotypic variation protein - fungus (Fusarium sporotrichioides) (fragment)  
C;Species: Fusarium sporotrichioides  
C;Date: 20-Feb-1995 #sequence\_revision 20-Feb-1995 #text\_change 20-Feb-1995  
C;Accession: PA0099  
R;Chow, L.P.; Fukaya, N.; Sugiura, Y.; Ueno, Y.; Tabuchi, K.; Tsugita, A.  
submitted to JIPID, October 1994  
A;Description: Two-dimensional polyacrylamide gel electrophoresis of Fusarium sporotrichioides  
A;Reference number: PA0051  
A;Accession: PA0099  
A;Molecule type: protein  
A;Residues: 1-15 <CHO>

Query Match 44.7%; Score 17; DB 2; Length 15;  
Best Local Similarity 66.7%; Pred. No. 3.3e+03;  
Matches 2; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

QY 4 WHF 6  
| |  
Db 5 WEF 7

RESULT 14  
T13818  
cytochrome oxidase subunit I - Atlantic hagfish mitochondrion (fragment)  
C;Species: mitochondrion Myxine glutinosa (Atlantic hagfish)  
C;Date: 20-Sep-1999 #sequence\_revision 20-Sep-1999 #text\_change 21-Jul-2000  
C;Accession: T13818  
R;Delarbre, C.; Barriol, V.; Tillier, S.; Janvier, P.; Gachelin, G.  
Mol. Biol. Evol. 14, 807-813, 1997  
A;Title: The main features of the craniate mitochondrial DNA between the ND1 and the C  
A;Reference number: Z17775; MUID:97398704; PMID:9254918  
A;Accession: T13818  
A;Status: preliminary; translated from GB/EMBL/DBJ

A;Molecule type: DNA  
A;Residues: 1-8 <DEL>  
A;Cross-references: EMBL:Y09527; NID:g2340019; PIDN:CAA70718.1; PID:g2340022  
C;Genetics:  
A;Genome: mitochondrion  
A;Note: COI  
C;Keywords: mitochondrion

Query Match 42.1%; Score 16; DB 2; Length 8;  
Best Local Similarity 66.7%; Pred. No. 2.8e+05;  
Matches 2; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

Qy 4 WHF 6  
| |  
Db 6 WFF 8

RESULT 15  
PQ0177  
neuromedin C - laughing frog  
C;Species: Rana ridibunda (laughing frog)  
C;Date: 23-Nov-1991 #sequence\_revision 23-Nov-1991 #text\_change 11-Jan-2000  
C;Accession: PQ0177  
R;Conlon, J.M.; O'Harte, F.; Vaudry, H.  
Biochem. Biophys. Res. Commun. 178, 526-530, 1991  
A;Title: Primary structures of the bombesin-like neuropeptides in frog brain show that b  
A;Reference number: PQ0177; MUID:91315477; PMID:1859413  
A;Accession: PQ0177  
A;Molecule type: protein  
A;Residues: 1-10 <CON>  
A;Experimental source: brain  
C;Superfamily: gastrin-releasing peptide  
C;Keywords: amidated carboxyl end  
F;10/Modified site: amidated carboxyl end (Met) #status predicted

Query Match 42.1%; Score 16; DB 2; Length 10;  
Best Local Similarity 40.0%; Pred. No. 3.4e+03;  
Matches 2; Conservative 0; Mismatches 3; Indels 0; Gaps 0;

Qy 1 WXXWH 5  
| |  
Db 4 WAVGH 8

Search completed: December 3, 2003, 11:54:09  
Job time : 12 secs

GenCore version 5.1.6  
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OM protein - protein search, using sw model

Run on: December 3, 2003, 11:44:55 ; Search time 7.33333 Seconds  
(without alignments)  
38.476 Million cell updates/sec

Title: US-09-912-414-11  
Perfect score: 38  
Sequence: 1 WXXWHF 6

Scoring table: BLOSUM62  
Gapop 10.0 , Gapext 0.5

Searched: 127863 seqs, 47026705 residues  
Total number of hits satisfying chosen parameters: 795

Minimum DB seq length: 0  
Maximum DB seq length: 15

Post-processing: Minimum Match 0%  
Maximum Match 100%  
Listing first 45 summaries

Database : SwissProt\_41:\*

Pred. No. is the number of results predicted by chance to have a  
score greater than or equal to the score of the result being printed,  
and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	DB ID	Description
1	23	60.5	9	1 LITR PHYRO	P08946 phyllomedusa
2	23	60.5	11	1 RANC_RANPI	P08951 rana pipien
3	22	57.9	9	1 LITO_LITAU	P08945 litoria aur
4	22	57.9	13	1 BOML_PSEGU	P42991 pseudophryn
5	21	55.3	10	1 LABA_JATMU	P13270 jatropha mu
6	19	50.0	9	1 COW_CONVE	P83047 conus ventr
7	16	42.1	10	1 GON2_CHEPR	P80678 chelyosoma
8	16	42.1	10	1 GRP_RANRI	P23260 rana ridibu
9	16	42.1	14	1 ALYT_ALYOB	P08944 alytes obst
10	15	39.5	15	1 RM12_YEAST	P36522 saccharomyc
11	14	36.8	10	1 FARP_MYTED	P42560 mytilus edu
12	14	36.8	11	1 CA22_LITCI	P82088 litoria cit
13	14	36.8	11	1 CA42_LITCI	P82092 litoria cit
14	14	36.8	11	1 MLG_THETS	P41989 theromyzon
15	14	36.8	13	1 CXA2_CONGE	P01520 conus geogr
16	14	36.8	13	1 MLA_ANOCA	P41589 anolis caro
17	14	36.8	13	1 MLA_CAMDR	P01198 camelus dro
18	14	36.8	15	1 AH2_PRUSE	P29260 prunus sero
19	14	36.8	15	1 DCMN_PSECH	P19917 pseudomonas
20	13	34.2	10	1 APE_CAPGI	P80474 capnocytoph
21	13	34.2	10	1 GON1_ALLMI	P37041 alligator m
22	13	34.2	10	1 GON2_CHICK	P37043 gallus gall
23	13	34.2	10	1 GON3_ONCKE	P20367 oncorhynch
24	13	34.2	12	1 UR2A_CATCO	P04558 catostomus
25	13	34.2	12	1 UR2B_CATCO	P04559 catostomus
26	13	34.2	12	1 UR2B_CYPCA	P04561 cyprinus ca
27	13	34.2	12	1 UR2_GILMI	P01147 gillichthys
28	13	34.2	12	1 UR2_POLSP	P81022 polyodon sp
29	13	34.2	12	1 UR2_SCYCA	P35490 scyliorhinu
30	13	34.2	15	1 UC16_MAIZE	P80622 zea mays (m
31	12	31.6	6	1 LOK1_LOCMI	P41491 locusta mig
32	12	31.6	8	1 LCK2_LEUMA	P21141 leucophaea
33	12	31.6	8	1 LCK5_LEUMA	P19987 leucophaea

34	12	31.6	8	1 LCK7_LEUMA	P19989 leucophaea
35	12	31.6	10	1 AEGL_AGRAE	P83465 agrocyebe ae
36	12	31.6	10	1 CA12_LITCI	P82086 litoria cit
37	12	31.6	10	1 CAER_LITXA	P56264 litoria xan
38	12	31.6	10	1 GONI_CHEPR	P80677 chelyosoma
39	12	31.6	10	1 HTF_TABAT	P14596 tabanus atr
40	12	31.6	11	1 RR2_CONAM	P42341 conopholis
41	12	31.6	15	1 RBS_PHYPA	P80657 physcomitre
42	11	28.9	4	1 OCP3_OCTMI	P58649 octopus min
43	11	28.9	5	1 BPP7_BOTIN	P30425 bothriops in
44	11	28.9	5	1 UF01_MOUSE	P38639 mus musculu
45	11	28.9	6	1 EI01_LITRU	P82096 litoria rub

ALIGNMENTS

RESULT 1

ID	LITR_PHYRO	STANDARD;	PRT;	9 AA.
AC	P08946;			
DT	01-NOV-1988 (Rel. 09, Created)			
DT	01-FEB-1994 (Rel. 28, Last sequence update)			
DT	15-SEP-2003 (Rel. 42, Last annotation update)			
DE	Rhodei-litorin.			
OS	Phyllomedusa rohdei (Rohde's leaf frog).			
OC	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;			
OC	Amphibia; Batrachia; Anura; Neobatrachia; Bufonoidea; Hylidae;			
OC	Phyllomedusinae; Phyllomedusa.			
OX	NCBI_TaxID=8394;			
RN	[1]			
RP	SEQUENCE.			
RC	TISSUE=Skin secretion;			
RX	MEDLINE=85127560; PubMed=3838283;			
RA	Barra D., Erspamer G.F., Simmaco M., Bossa F., Melchiorri P.,			
RA	Erspamer V.;			
RT	"Rohdei-litorin: a new peptide from the skin of Phyllomedusa rohdei.";			
RL	FEBS Lett. 182:53-56(1985).			
CC	- - SUBCELLULAR LOCATION: Secreted.			
CC	- - TISSUE SPECIFICITY: Skin.			
CC	- - SIMILARITY: BELONGS TO THE BOMBESIN/NEUROMEDIN B/RANATENSIN FAMILY.			
CC	PIR; S07241; S07241.			
DR	InterPro; IPR000874; Bombesin.			
DR	Pfam; PF02044; Bombesin; 1.			
DR	PROSITE; PS00257; BOMBESIN; 1.			
KW	Amphibian defense peptide; Bombesin family; Amidation;			
KW	Pyrolidone carboxylic acid.			
FT	MOD_RES 1 1 PYRROLIDONE CARBOXYLIC ACID.			
FT	MOD_RES 9 9 AMIDATION.			
SQ	SEQUENCE 9 AA; 1090 MW; 4ECCC1E861ADC377 CRC64;			

Query Match 60.5%; Score 23; DB 1; Length 9;  
Best Local Similarity 50.0%; Pred. No. 1.3e+05;  
Matches 3; Conservative 0; Mismatches 3; Indels 0; Gaps 0;

QY	1	WXXWHF 6
Db	3	WATGHF 8

RESULT 2

ID	RANC_RANPI	STANDARD;	PRT;	11 AA.
AC	P08951;			
DT	01-NOV-1988 (Rel. 09, Created)			
DT	01-NOV-1988 (Rel. 09, Last sequence update)			
DT	15-SEP-2003 (Rel. 42, Last annotation update)			
DE	Ranatensin-C.			
OS	Rana pipiens (Northern leopard frog).			
OC	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;			
OC	Amphibia; Batrachia; Anura; Neobatrachia; Ranioidea; Rana.			
OX	NCBI_TaxID=8404;			

RN [1]  
RP SEQUENCE.  
RC TISSUE=Skin secretion;  
RX MEDLINE=84131098; PubMed=6141890;  
RA Nakajima T.;  
RL Unpublished results, cited by:  
RL Erspamer V., Erspamer G.F., Mazzanti G., Endean R.;  
RL Comp. Biochem. Physiol. 77C:99-108(1984).  
CC -!- SUBCELLULAR LOCATION: Secreted.  
CC -!- TISSUE SPECIFICITY: Skin.  
CC -!- SIMILARITY: BELONGS TO THE BOMBESIN/NEUROMEDIN B/RANATENSIN FAMILY.  
CC InterPro; IPR000874; Bombesin.  
DR Pfam; PF02044; Bombesin; 1.  
DR PROSITE; PS00257; BOMBESIN; 1.  
KW Amphibian defense peptide; Bombesin family; Amidation.  
FT MOD\_RES 11 11 AMIDATION.  
SQ SEQUENCE 11 AA; 1304 MW; D6C9885A61ADC366 CRC64;  
  
Query Match 60.5%; Score 23; DB 1; Length 11;  
Best Local Similarity 50.0%; Pred. No. 94;  
Matches 3; Conservative 0; Mismatches 3; Indels 0; Gaps 0;  
  
QY 1 WXXWHF 6  
Db 5 WATGHF 10  
  
RESULT 3  
LITO LITAU STANDARD; PRT; 9 AA.  
AC P08945;  
DT 01-NOV-1988 (Rel. 09, Created)  
DT 01-FEB-1994 (Rel. 28, Last sequence update)  
DT 15-SEP-2003 (Rel. 42, Last annotation update)  
DE Litorin.  
OS Litoria aurea (Green and golden bell frog).  
OC Eukaryota; Metazoa; Chordata; Vertebrata; Euteleostomi;  
OC Amphibia; Batrachia; Anura; Neobatrachia; Bufonoidea; Hylidae;  
OC Pelodyadinae; Litoria.  
OX NCBI\_TaxID=8371;  
RN [1]  
RP SEQUENCE.  
RC TISSUE=Skin secretion;  
RX MEDLINE=75187011; PubMed=1140241;  
RA Anastasi A., Erspamer V., Endean R.;  
RT "Amino acid composition and sequence of litorin, a bombesin-like nonapeptide from the skin of the Australian leptodactylid frog Litoria aurea.";  
RT Litoria aurea.";  
RL Experientia 31:510-511(1975).  
RN [2]  
RP SEQUENCE (METHYLATED VARIANT).  
RC TISSUE=Skin secretion;  
RX MEDLINE=78003546; PubMed=908397;  
RA Anastasi A., Montecucchi P.C., Angelucci F., Erspamer V., Endean R.;  
RT "Glu(OMe)3-litorin, the second bombesin-like peptide occurring in methanol extracts of the skin of the Australian frog Litoria aurea.";  
RL Experientia 33:1289-1289(1977).  
CC -!- SUBCELLULAR LOCATION: Secreted.  
CC -!- TISSUE SPECIFICITY: Skin.  
CC -!- SIMILARITY: BELONGS TO THE BOMBESIN/NEUROMEDIN B/RANATENSIN FAMILY.  
CC PIR; S07204; S07204.  
DR InterPro; IPR000874; Bombesin.  
DR Pfam; PF02044; Bombesin; 1.  
DR PROSITE; PS00257; BOMBESIN; 1.  
KW Amphibian defense peptide; Bombesin family; Amidation; Methylation;  
KW Pyrrolidone carboxylic acid.  
FT MOD\_RES 1 1 PYRROLIDONE CARBOXYLIC ACID.  
FT MOD\_RES 2 2 METHYLATION (PARTIAL).  
FT MOD\_RES 9 9 AMIDATION.  
SQ SEQUENCE 9 AA; 1103 MW; D7CCC1E862CDC366 CRC64;

Query Match 57.9%; Score 22; DB 1; Length 9;  
Best Local Similarity 50.0%; Pred. No. 1.3e+05;  
Matches 3; Conservative 0; Mismatches 3; Indels 0; Gaps 0;  
  
QY 1 WXXWHF 6  
Db 3 WAVGHF 8  
  
RESULT 4  
BOML\_PSEGU STANDARD; PRT; 13 AA.  
AC P42991;  
DT 01-NOV-1995 (Rel. 32, Created)  
DT 01-NOV-1995 (Rel. 32, Last sequence update)  
DT 15-SEP-2003 (Rel. 42, Last annotation update)  
DE Bombesin-like peptide L (PG-L).  
OS Pseudophryne guentheri (Guenther's toadlet).  
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
OC Amphibia; Batrachia; Anura; Neobatrachia; Bufonoidea; Myobatrachidae;  
OC Myobatrachinae; Pseudophryne.  
OX NCBI\_TaxID=30349;  
RN [1]  
RP SEQUENCE.  
RC TISSUE=Skin secretion;  
RX MEDLINE=90287814; PubMed=2356157;  
RA Simmaco M., Severini C., de Biase D., Barra D., Bossa F.,  
RA Roberts J.D., Melchiorri P., Erspamer V.;  
RT "Six novel tachykinin- and bombesin-related peptides from the skin of the Australian frog Pseudophryne guntheri.";  
RL Peptides 11:299-304(1990).  
CC -!- SUBCELLULAR LOCATION: Secreted.  
CC -!- TISSUE SPECIFICITY: Skin.  
CC -!- SIMILARITY: BELONGS TO THE BOMBESIN/NEUROMEDIN B/RANATENSIN FAMILY.  
CC PIR; A60409; A60409.  
DR InterPro; IPR000874; Bombesin.  
DR Pfam; PF02044; Bombesin; 1.  
DR PROSITE; PS00257; BOMBESIN; 1.  
KW Amphibian defense peptide; Bombesin family; Amidation;  
KW Pyrrolidone carboxylic acid.  
FT MOD\_RES 1 1 PYRROLIDONE CARBOXYLIC ACID.  
FT MOD\_RES 13 13 AMIDATION.  
SQ SEQUENCE 13 AA; 1372 MW; D6DE0D24BD98C366 CRC64;  
  
Query Match 57.9%; Score 22; DB 1; Length 13;  
Best Local Similarity 50.0%; Pred. No. 1.6e+02;  
Matches 3; Conservative 0; Mismatches 3; Indels 0; Gaps 0;

QY 1 WXXWHF 6  
Db 7 WAVGHF 12  
  
RESULT 5  
LABA\_JATMU STANDARD; PRT; 10 AA.  
AC P13270;  
DT 01-JAN-1990 (Rel. 13, Created)  
DT 01-JAN-1990 (Rel. 13, Last sequence update)  
DT 28-FEB-2003 (Rel. 41, Last annotation update)  
DE Labaditin.  
OS Jatropha multifida (Physic nut).  
OC Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;  
OC Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; Rosidae;  
OC eucosids I; Malpighiales; Euphorbiaceae; Jatropha.  
OX NCBI\_TaxID=3996;  
RN [1]  
RP SEQUENCE.  
RC TISSUE=Latex;  
RA Kosasi S., van der Sluis W.G., Boelens R., T'Hart L.A., Labadie R.P.;  
RT "Labaditin, a novel cyclic decapeptide from the latex of Jatropha multifida L. (Euphorbiaceae). Isolation and sequence determination



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RT by means of two-dimensional NMR." ;
RL FEBS Lett. 256:91-96(1989).
CC -!- FUNCTION: LABADITIN IS AN ACTIVE PEPTIDE WHICH INHIBITS THE
CC CLASSICAL PATHWAY OF COMPLEMENT ACTIVATION IN VITRO. ACTIVITY
CC SEEMS TO BE BASED ON AN INTERACTION WITH C1.
CC -!- PTM: This is a cyclic peptide.
CC -!- DISEASE: LATEX OF THIS PLANT IS USED IN FOLKLORIC MEDICINE FOR
CC TREATMENT OF INFECTED WOUNDS, SKINS INFECTIONS AND SCABIES.
SQ SEQUENCE 10 AA; 1089 MW; D98AAD6362D1B362 CRC64;

Query Match 55.3%; Score 21; DB 1; Length 10;
Best Local Similarity 50.0%; Pred. No. 1.9e+02;
Matches 2; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

QY 1 WXXW 4
| |
Db 4 WTVW 7

RESULT 6
COW_CONVE STANDARD; PRT; 9 AA.
AC P83047;
DT 16-OCT-2001 (Rel. 40, Created)
DT 16-OCT-2001 (Rel. 40, Last sequence update)
DT 28-FEB-2003 (Rel. 41, Last annotation update)
DE Contryphan-Vn.
OS Conus ventricosus (Mediterranean cone).
OC Eukaryota; Metazoa; Mollusca; Gastropoda; Orthogastropoda;
OC Apogastropoda; Caenogastropoda; Sorbeoconcha; Hypsogastropoda;
OC Neogastropoda; Conoidea; Conidae; Conus.
OX NCBI_TaxID=117992;
RN [1]
RP SEQUENCE, SYNTHESIS, AND MASS SPECTROMETRY.
RC TISSUE=Venom;
RX MEDLINE=21547785; PubMed=11688995;
RA Massilia G.R., Schinina M.E., Ascenzi P., Polticelli F.;
RT "Contryphan-Vn: a novel peptide from the venom of the Mediterranean
RT snail Conus ventricosus.";
RL Biochem. Biophys. Res. Commun. 288:908-913(2001).
CC -!- SUBCELLULAR LOCATION: Secreted.
CC -!- TISSUE SPECIFICITY: Expressed by the venom duct.
CC -!- MASS SPECTROMETRY: MW=1088.6; METHOD=MALDI.
CC -!- SIMILARITY: BELONGS TO THE CONTRYPHAN FAMILY.
KW Toxin; Amidation; D-amino acid.
FT DISULFID 3 9
FT MOD_RES 5 5 D-TRYPTOPHAN.
FT MOD_RES 9 9 AMIDATION.
FT SEQUENCE 9 AA; 1091 MW; 8D38676323676EBA CRC64;

Query Match 50.0%; Score 19; DB 1; Length 9;
Best Local Similarity 50.0%; Pred. No. 1.3e+05;
Matches 2; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

QY 1 WXXW 4
| |
Db 5 WKPW 8

RESULT 7
GON2_CHEPR STANDARD; PRT; 10 AA.
AC P80678;
DT 01-NOV-1997 (Rel. 35, Created)
DT 01-NOV-1997 (Rel. 35, Last sequence update)
DT 28-FEB-2003 (Rel. 41, Last annotation update)
DE Gonadoliberin II (Gonadotropin-releasing hormone II) (GnRH-II)
DE (Luliberin II).
OS Chelyosoma productum.
OC Eukaryota; Metazoa; Chordata; Urochordata; Ascidiacea; Enterogona;
OC Phlebobranchia; Corellidae; Chelyosoma.
OX NCBI_TaxID=71177;
RN [1]
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RP SEQUENCE.
RX MEDLINE=96413669; PubMed=8816823;
RA Powell J.F.F., Reska-Skinner S.M., Prakash M.O., Fischer W.H.,
RA Park M., Rivier J.E., Craig A.G., Mackie G.O., Sherwood N.M.;
RT "Two new forms of gonadotropin-releasing hormone in a protochordate
RT and the evolutionary implications.";
RL Proc. Natl. Acad. Sci. U.S.A. 93:10461-10464(1996).
CC -!- FUNCTION: Stimulates the secretion of gonadotropins; it stimulates
CC the secretion of both luteinizing and follicle-stimulating
CC hormones.
CC -!- SUBUNIT: Homodimer; disulfide-linked.
CC -!- SUBCELLULAR LOCATION: Secreted.
CC -!- TISSUE SPECIFICITY: GNRH NEURONS LIE WITHIN BLOOD SINUSES CLOSE TO
CC THE GONODUCTS AND GONADS IN BOTH JUVENILES AND ADULTS, IMPLYING
CC THAT THE NEUROPEPTIDE IS RELEASED INTO THE BLOODSTREAM.
CC -!- MASS SPECTROMETRY: MW=1117.52; METHOD=MALDI.
CC -!- SIMILARITY: Belongs to the GNRH family.
DR InterPro; IPR002012; GNRH.
DR PROSITE; PS00473; GNRH; 1.
KW Hormone; Amidation; Pyrrolidone carboxylic acid.
FT MOD_RES 1 1 PYRROLIDONE CARBOXYLIC ACID.
FT DISULFID 6 6 INTERCHAIN.
FT MOD_RES 10 10 AMIDATION (BY SIMILARITY).
SQ SEQUENCE 10 AA; 1135 MW; 284B38D1EEB735A3 CRC64;

Query Match 42.1%; Score 16; DB 1; Length 10;
Best Local Similarity 40.0%; Pred. No. 1.3e+03;
Matches 2; Conservative 0; Mismatches 3; Indels 0; Gaps 0;

QY 1 WXXWH 5
| |
Db 3 WSLCH 7

RESULT 8
GRP_RANRI STANDARD; PRT; 10 AA.
ID_P23260;
AC P23260;
DT 01-NOV-1991 (Rel. 20, Created)
DT 01-NOV-1991 (Rel. 20, Last sequence update)
DT 28-FEB-2003 (Rel. 41, Last annotation update)
DE Neuromedin C.
OS Rana ridibunda (laughing frog) (Marsh frog).
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Amphibia; Batrachia; Anura; Neobatrachia; Ranioidea; Rana.
OX NCBI_TaxID=8406;
RN [1]
RP SEQUENCE.
RC TISSUE=Brain;
RX MEDLINE=91315477; PubMed=1859413;
RA Conlon J.M., O'Harte F., Vaudry H.;
RT "Primary structures of the bombesin-like neuropeptides in frog brain
RT show that bombesin is not the amphibian gastrin-releasing peptide.";
RL Biochem. Biophys. Res. Commun. 178:526-530(1991).
CC -!- SUBCELLULAR LOCATION: Secreted.
CC -!- SIMILARITY: BELONGS TO THE BOMBESIN/NEUROMEDIN B/RANATENSIN
CC FAMILY.
DR PIR; PQ0177; PQ0177.
DR InterPro; IPR000874; Bombesin.
DR Pfam; PF02044; Bombesin; 1.
DR PROSITE; PS00257; BOMBESIN; 1.
KW Bombesin family; Amidation.
FT MOD_RES 10 10 AMIDATION.
FT SEQUENCE 10 AA; 1094 MW; F81FBA862CDC371 CRC64;

Query Match 42.1%; Score 16; DB 1; Length 10;
Best Local Similarity 40.0%; Pred. No. 1.3e+03;
Matches 2; Conservative 0; Mismatches 3; Indels 0; Gaps 0;

QY 1 WXXWH 5
| |
Db 4 WAVGH 8
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RESULT 9
ALYT_ALYOB
ID ALYT_ALYOB STANDARD; PRT; 14 AA.
AC P08944;
DT 01-NOV-1988 (Rel. 09, Created)
DT 01-FEB-1994 (Rel. 28, Last sequence update)
DT 15-SEP-2003 (Rel. 42, Last annotation update)
DE Alytesin.
OS Alytes obstetricans (Midwife toad).
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Amphibia; Batrachia; Anura; Archeobatrachia; Discoglossidae; Alytes.
OX NCBI_TaxID=8443;
RN [1]
RP SEQUENCE.
RC TISSUE=Skin secretion;
RX MEDLINE=84131098; PubMed=6141890;
RA Erspamer V., Erspamer G.F., Mazzanti G., Endean R.;
RT "Active peptides in the skins of one hundred amphibian species from
Australia and Papua New Guinea.";
RL Comp. Biochem. Physiol. 77C:99-108(1984).
CC -!- SUBCELLULAR LOCATION: Secreted.
CC -!- TISSUE SPECIFICITY: Skin.
CC -!- SIMILARITY: BELONGS TO THE BOMBESIN/NEUROMEDIN B/RANATENSIN
FAMILY.
DR InterPro; IPR000874; Bombesin.
DR Pfam; PF02044; Bombesin; 1.
DR PROSITE; PS00257; BOMBESIN; 1.
KW Amphibian defense peptide; Bombesin family; Amidation;
KW Pyrrolidone carboxylic acid.
FT MOD_RES 1 1 PYRROLIDONE CARBOXYLIC ACID.
FT MOD_RES 14 14 AMIDATION.
SQ SEQUENCE 14 AA; 1554 MW; D3C4E4D3AF129666 CRC64;

Query Match 42.1%; Score 16; DB 1; Length 14;
Best Local Similarity 40.0%; Pred. No. 1.7e+03;
Matches 2; Conservative 0; Mismatches 3; Indels 0; Gaps 0;

QY 1 WXXWH 5
Db 8 WAVGH 12

RESULT 10
RM12_YEAST
ID RM12_YEAST STANDARD; PRT; 15 AA.
AC P36522;
DT 01-JUN-1994 (Rel. 29, Created)
DT 01-JUN-1994 (Rel. 29, Last sequence update)
DT 01-JUN-1994 (Rel. 29, Last annotation update)
DE Mitochondrial 60S ribosomal protein L12 (YmL12) (Fragment).
GN MRPL12.
OS Saccharomyces cerevisiae (Baker's yeast).
OC Eukaryota; Fungi; Ascomycota; Saccharomycotina; Saccharomycetes;
OC Saccharomycetales; Saccharomycetaceae; Saccharomycetes.
OX NCBI_TaxID=4932;
RN [1]
RP SEQUENCE.
RX MEDLINE=91285106; PubMed=2060626;
RA Grohmann L., Graack H.-R., Kruft V., Choli T., Goldschmidt-Reisin S.,
RA Kitakawa M.;
RT "Extended N-terminal sequencing of proteins of the large ribosomal
subunit from yeast mitochondria.";
RL FEBS Lett. 284:51-56(1991).
DR SGD; L0002687; MRPL12.
KW Ribosomal protein; Mitochondrion.
FT NON_TER 15 15
SQ SEQUENCE 15 AA; 1851 MW; 74BCD9FEDDDDB3900 CRC64;

Query Match 39.5%; Score 15; DB 1; Length 15;
Best Local Similarity 50.0%; Pred. No. 2.7e+03;
Matches 3; Conservative 1; Mismatches 2; Indels 0; Gaps 0;
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QY 1 WXXWHF 6
Db 1 WXDGYF 6

RESULT 11
FARP_MYTED
ID FARP_MYTED STANDARD; PRT; 10 AA.
AC P42560;
DT 01-NOV-1995 (Rel. 32, Created)
DT 01-NOV-1995 (Rel. 32, Last sequence update)
DT 30-MAY-2000 (Rel. 39, Last annotation update)
DE FMRFamide-like neuropeptide ALAGDHFFRF-amide.
OS Mytilus edulis (Blue mussel).
OC Eukaryota; Metazoa; Mollusca; Bivalvia; Pteriomorpha; Mytiloida;
OC Mytiloidea; Mytilidae; Mytilus.
OX NCBI_TaxID=6550;
RN [1]
RP SEQUENCE.
RX MEDLINE=93047883; PubMed=1358534;
RA Walker R.J.;
RT "Neuroactive peptides with an RFamide or Famide carboxyl terminal.";
RL Comp. Biochem. Physiol. 102C:213-222(1992).
CC -!- SIMILARITY: BELONGS TO THE FARP (FMRFAMIDE RELATED PEPTIDE)
FAMILY.
CC PIR; A58365; A58365.
KW Neuropeptide; Amidation.
FT MOD_RES 10 10 AMIDATION.
SQ SEQUENCE 10 AA; 1180 MW; C2F80CC9C1EAA87D CRC64;

Query Match 36.8%; Score 14; DB 1; Length 10;
Best Local Similarity 100.0%; Pred. No. 2.7e+03;
Matches 2; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 5 HF 6
Db 6 HF 7

RESULT 12
CA22_LITCI
ID CA22_LITCI STANDARD; PRT; 11 AA.
AC P82088;
DT 16-OCT-2001 (Rel. 40, Created)
DT 16-OCT-2001 (Rel. 40, Last sequence update)
DT 15-SEP-2003 (Rel. 42, Last annotation update)
DE Caerulein 2.2/2.2Y4.
OS Litoria citropa (Australian blue mountains tree frog).
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Amphibia; Batrachia; Anura; Neobatrachia; Bufonoidea; Hylidae;
OC Pelodyadinae; Litoria.
OX NCBI_TaxID=94770;
RN [1]
RP SEQUENCE, AND MASS SPECTROMETRY.
RC TISSUE=Skin secretion;
RX MEDLINE=20057701; PubMed=10589099;
RA Wabnitz P.A., Bowie J.H., Tyler M.J.;
RT "Caerulein-like peptides from the skin glands of the Australian blue
mountains tree frog Litoria citropa. Part 1. Sequence determination
using electrospray mass spectrometry.";
RL Rapid Commun. Mass Spectrom. 13:2498-2502(1999).
CC -!- FUNCTION: HYPOTENSIVE NEUROPEPTIDE (PROBABLE).
CC -!- SUBCELLULAR LOCATION: Secreted.
CC -!- TISSUE SPECIFICITY: Skin dorsal glands.
CC -!- PTM: Isoform 2.2Y4 differs from isoform 2.2 in not being
sulfated.
CC -!- MASS SPECTROMETRY: MW=1388; METHOD=Electrospray.
CC -!- SIMILARITY: BELONGS TO THE GASTRIN/CHOLECYSTOKININ FAMILY.
DR InterPro; IPR001651; Gastrin.
DR PROSITE; PS00259; GASTRIN; FALSE_NEG.
KW Amphibian defense peptide; Hypotensive agent; Amidation; Sulfation;
KW Pyrrolidone carboxylic acid.
FT MOD_RES 1 1 PYRROLIDONE CARBOXYLIC ACID.
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FT  MOD RES      4      4      SULFATION.
FT  MOD_RES      11     11     AMIDATION.
SQ  SEQUENCE     11 AA; 1328 MW; 10DAB894EDD861BB CRC64;

Query Match      36.8%; Score 14; DB 1; Length 11;
Best Local Similarity 100.0%; Pred. No. 3e+03;
Matches 2; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY      5 HF 6
Db      8 HF 9

RESULT 13
CA42_LITCI      STANDARD; PRT; 11 AA.
AC  P82092;
DT  16-OCT-2001 (Rel. 40, Created)
DT  16-OCT-2001 (Rel. 40, Last sequence update)
DT  15-SEP-2003 (Rel. 42, Last annotation update)
DE  Caerulein 4.2/4.2Y4.
OS  Litoria citropa (Australian blue mountains tree frog).
OC  Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC  Amphibia; Batrachia; Anura; Neobatrachia; Bufonoidea; Hylidae;
OC  Pelodyadinae; Litoria.
OX  NCBI_TaxID=94770;
RN  [1]
RP  SEQUENCE, AND MASS SPECTROMETRY.
RC  TISSUE=Skin secretion;
RX  MEDLINE=20057701; PubMed=10589099;
RA  Wabnitz P.A., Bowie J.H., Tyler M.J.;
RT  "Caerulein-like peptides from the skin glands of the Australian blue
RT  mountains tree frog Litoria citropa. Part 1. Sequence determination
RT  using electrospray mass spectrometry.";
RL  Rapid Commun. Mass Spectrom. 13:2498-2502(1999).
CC  -!- FUNCTION: HYPOTENSIVE NEUROPEPTIDE (PROBABLE).
CC  -!- SUBCELLULAR LOCATION: Secreted.
CC  -!- TISSUE SPECIFICITY: Skin dorsal glands.
CC  -!- PTM: Isoform 4.2Y4 differs from isoform 4.2 in not being
CC  sulfated.
CC  -!- MASS SPECTROMETRY: MW=1404; METHOD=Electrospray.
CC  -!- SIMILARITY: BELONGS TO THE GASTRIN/CHOLECYSTOKININ FAMILY.
DR  InterPro; IPR001651; Gastrin.
DR  PROSITE; PS00259; GASTRIN; FALSE_NEG.
KW  Amphibian defense peptide; Hypotensive agent; Amidation; Sulfation;
KW  Pyrrolidone carboxylic acid.
FT  MOD_RES      1      1      PYRROLIDONE CARBOXYLIC ACID.
FT  MOD_RES      4      4      SULFATION.
FT  MOD_RES      11     11     AMIDATION.
SQ  SEQUENCE     11 AA; 1344 MW; 10DAB894F5B861BB CRC64;

Query Match      36.8%; Score 14; DB 1; Length 11;
Best Local Similarity 100.0%; Pred. No. 3e+03;
Matches 2; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY      5 HF 6
Db      8 HF 9

RESULT 14
MLG_THETS
ID  MLG THETS      STANDARD; PRT; 11 AA.
AC  P41989;
DT  01-NOV-1995 (Rel. 32, Created)
DT  01-NOV-1995 (Rel. 32, Last sequence update)
DT  16-OCT-2001 (Rel. 40, Last annotation update)
DE  Melanotropin gamma (Gamma-melanocyte stimulating hormone) (Gamma-MSH).
OS  Theromyzon tessulatum (Leech).
OC  Eukaryota; Metazoa; Annelida; Clitellata; Hirudinida; Hirudinea;
OC  Rhynchobdellida; Glossiphoniidae; Theromyzon.
OX  NCBI_TaxID=13286;
RN  [1]
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RP  SEQUENCE.
RC  TISSUE=Brain;
RX  MEDLINE=94298944; PubMed=8026574;
RA  Salzet M., Wattez C., Bulet P., Malecha J.;
RT  "Isolation and structural characterization of a novel peptide related
RT  to gamma-melanocyte stimulating hormone from the brain of the leech
RT  Theromyzon tessulatum.";
RL  FEBS Lett. 348:102-106(1994).
CC  -!- SIMILARITY: BELONGS TO THE POMC FAMILY.
DR  PIR; S45698; S45698.
KW  Hormone; Amidation.
FT  MOD_RES      11     11     AMIDATION.
SQ  SEQUENCE     11 AA; 1486 MW; 2DB8FACE6409C1E8 CRC64;

Query Match      36.8%; Score 14; DB 1; Length 11;
Best Local Similarity 100.0%; Pred. No. 3e+03;
Matches 2; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY      5 HF 6
Db      5 HF 6

RESULT 15
CX42_CONGE      STANDARD; PRT; 13 AA.
AC  P01520;
DT  21-JUL-1986 (Rel. 01, Created)
DT  21-JUL-1986 (Rel. 01, Last sequence update)
DT  28-FEB-2003 (Rel. 41, Last annotation update)
DE  Alpha-conotoxin GII.
OS  Conus geographus (Geography cone).
OC  Eukaryota; Metazoa; Mollusca; Gastropoda; Orthogastropoda;
OC  Apogastropoda; Caenogastropoda; Sorbeoconcha; Hypsogastropoda;
OC  Neogastropoda; Conoidea; Conidae; Conus.
OX  NCBI_TaxID=6491;
RN  [1]
RP  SEQUENCE.
RX  MEDLINE=81191854; PubMed=7014556;
RA  Gray W.R., Luque A., Olivera B.M., Barrett J., Cruz L.J.;
RT  "Peptide toxins from Conus geographus venom.";
RL  J. Biol. Chem. 256:4734-4740(1981).
RN  [2]
RP  DISULFIDE BONDS.
RX  MEDLINE=84280842; PubMed=6466616;
RA  Gray W.R., Luque F.A., Galyean R., Atherton E., Sheppard R.C.,
RA  Stone B.L., Reyes A., Alford J., McIntosh M., Olivera B.M.,
RA  Cruz L.J., Rivier J.;
RT  "Conotoxin GI: disulfide bridges, synthesis, and preparation of
RT  iodinated derivatives.";
RL  Biochemistry 23:2796-2802(1984).
CC  -!- FUNCTION: Alpha-conotoxins act on postsynaptic membranes, they
CC  bind to the nicotinic acetylcholine receptors (nAChR) and thus
CC  inhibit them.
CC  -!- SUBCELLULAR LOCATION: Secreted.
CC  -!- TISSUE SPECIFICITY: Expressed by the venom duct.
CC  -!- SIMILARITY: BELONGS TO THE A-SUPERFAMILY OF CONOTOXINS. ALPHA-TYPE
CC  FAMILY.
DR  PIR; A01783; NTKN2G.
DR  HSSP; P56973; 1B45.
KW  Postsynaptic neurotoxin; Neurotoxin; Toxin;
KW  Acetylcholine receptor inhibitor; Amidation.
FT  DISULFID      2      7
FT  DISULFID      3     13
FT  MOD_RES      13     13     AMIDATION.
SQ  SEQUENCE     13 AA; 1422 MW; DEE831C39297EBD CRC64;

Query Match      36.8%; Score 14; DB 1; Length 13;
Best Local Similarity 100.0%; Pred. No. 3.4e+03;
Matches 2; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY      5 HF 6
Db      5 HF 6
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Db 10 HF 11

Search completed: December 3, 2003, 11:51:52  
Job time : 8.33333 secs

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